

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208660_at	7.58E-04	gb:BC000105.1 /DEF=Homo sapiens, Similar to CG14740 gene product, clone MGC:2503, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG14740 gene product /DB_XREF=gi:12652712 /UG=Hs.239760 citrate synthase /FL=gb:BC000105.1 gb:AF047042.1 gb:NM_004077.1	BC000105		NP_004068
208661_s_at	0.018693	Consensus includes gb:AW510696 /FEA=EST /DB_XREF=gi:7148774 /DB_XREF=est:hc89f12.x1 /CLONE=IMAGE:2907215 /UG=Hs.118174 tetratricopeptide repeat domain 3 /FL=gb:D84294.1	D84294		NP_003307
208662_s_at	1.26E-04	Consensus includes gb:AI885338 /FEA=EST /DB_XREF=gi:5590502 /DB_XREF=est:wl92e09.x1 /CLONE=IMAGE:2432392 /UG=Hs.118174 tetratricopeptide repeat domain 3 /FL=gb:D84294.1	D84294		NP_003307
208663_s_at	0.027792	Consensus includes gb:AI652848 /FEA=EST /DB_XREF=gi:4736827 /DB_XREF=est:wb40a04.x1 /CLONE=IMAGE:2308110 /UG=Hs.118174 tetratricopeptide repeat domain 3 /FL=gb:D84294.1	D84294		NP_003307
208668_x_at	1.26E-04	gb:BC003689.1 /DEF=Homo sapiens, high-mobility group (nonhistone chromosomal) protein 17, clone MGC:5301, mRNA, complete cds. /FEA=mRNA /PROD=high-mobility group (nonhistone chromosomal)protein 17 /DB_XREF=gi:13277559 /UG=Hs.181163 high-mobility group (nonhistone chromosomal) protein 17 /FL=gb:BC003689.1 gb:M12623.1	BC003689		NP_005508

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208669_s_at	0.013223	gb:AF109873.1 /DEF=Homo sapiens retinoblastoma protein-associated protein mRNA, complete cds. /FEA=mRNA /PROD=retinoblastoma protein-associated protein /DB_XREF=gi:11415000 /UG=Hs.75847 CREBBPEP300 inhibitory protein 1 /FL=gb:AF109873.1 gb:AF274947.1 gb:AF274951.1 gb:AF092135.1 gb:NM_014335.1	AF109873		NP_055150
208671_at	1.26E-04	gb:AF164794.1 /DEF=Homo sapiens Diff33 protein homolog mRNA, complete cds. /FEA=mRNA /PROD=Diff33 protein homolog /DB_XREF=gi:8895090 /UG=Hs.146668 KIAA1253 protein /FL=gb:AF164794.1	AF164794		NP_065806
208674_x_at	1.26E-04	gb:BC002594.1 /DEF=Homo sapiens, dolichyl-diphosphooligosaccharide-protein glycosyltransferase, clone MGC:2191, mRNA, complete cds. /FEA=mRNA /PROD=dolichyl-diphosphooligosaccharide-proteinglycosyltransferase /DB_XREF=gi:12803530 /UG=Hs.34789 dolichyl-diphosphooligosaccharide-protein glycosyltransferase /FL=gb:BC002594.1 gb:D29643.1 gb:NM_005216.1	BC002594		NP_005207
208678_at	0.00161	gb:BC004443.1 /DEF=Homo sapiens, ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 31kD, clone MGC:4040, mRNA, complete cds. /FEA=mRNA /PROD=ATPase, H ⁺ transporting, lysosomal (vacuolarproton pump) 31kD /DB_XREF=gi:13325247 /UG=Hs.77805 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 31kD /FL=gb:BC004443.1 gb:NM_001696.1	BC004443		NP_001687

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208679_s_at	7.58E-04	gb:AF279893.1 /DEF=Homo sapiens PNAS-139 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-139 /DB_XREF=gi:12751114 /UG=Hs.83583 actin related protein 23 complex, subunit 2 (34 kD) /FL=gb:BC000590.1 gb:AF279893.1	AF279893		NP_690601
208684_at	0.001116	gb:U24105.1 /DEF=Homo sapiens coatomer protein (COPA) mRNA, complete cds. /FEA=mRNA /GEN=COPA /PROD=coatomer protein /DB_XREF=gi:1638873 /UG=Hs.75887 coatomer protein complex, subunit alpha /FL=gb:U24105.1 gb:NM_004371.2	U24105		NP_004362
208685_x_at	0.001054	bromodomain containing 2	AA902767	Hs.75243	NP_005095
208689_s_at	0.033381	gb:BC003560.1 /DEF=Homo sapiens, ribophorin II, clone MGC:1817, mRNA, complete cds. /FEA=mRNA /PROD=ribophorin II /DB_XREF=gi:13097707 /UG=Hs.75722 ribophorin II /FL=gb:BC003560.1 gb:NM_002951.1	BC003560		NP_002942
208691_at	0.001116	gb:BC001188.1 /DEF=Homo sapiens, transferrin receptor (p90, CD71), clone MGC:3151, mRNA, complete cds. /FEA=mRNA /PROD=transferrin receptor (p90, CD71) /DB_XREF=gi:12654696 /UG=Hs.77356 transferrin receptor (p90, CD71) /FL=gb:BC001188.1 gb:M11507.1	BC001188		NP_003225
208692_at	0.013223	gb:U14990.1 /DEF=Human XP1PO ribosomal protein S3 (rpS3) mRNA, complete cds. /FEA=mRNA /GEN=rpS3 /PROD=ribosomal protein S3 /DB_XREF=gi:555940 /UG=Hs.252259 ribosomal protein S3 /FL=gb:BC003137.1 gb:NM_001005.1 gb:U14990.1 gb:U14991.1 gb:U14992.1	U14990		NP_000996

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208696_at	1.26E-04	gb:AF275798.1 /DEF=Homo sapiens PNAS-102 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-102 /DB_XREF=gi:10834761 /UG=Hs.1600 chaperonin containing TCP1, subunit 5 (epsilon) /FL=gb:AF275798.1	AF275798		NP_036205
208702_x_at	0.003171	Consensus includes gb:AI525212 /FEA=EST /DB_XREF=gi:4439347 /DB_XREF=est:pt1.1-2.A08.r /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1	BC000373		NP_001633
208703_s_at	0.026891	Consensus includes gb:BG427393 /FEA=EST /DB_XREF=gi:13333995 /DB_XREF=est:602499110F1 /CLONE=IMAGE:4612562 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1	BC000373		NP_001633
208704_x_at	2.05E-04	gb:BC000373.1 /DEF=Homo sapiens, Similar to amyloid beta (A4) precursor-like protein 2, clone MGC:8371, mRNA, complete cds. /FEA=mRNA /PROD=Similar to amyloid beta (A4) precursor-likeprotein 2 /DB_XREF=gi:12653210 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1	BC000373		NP_001633
208706_s_at	0.041055	Consensus includes gb:AK026933.1 /DEF=Homo sapiens cDNA: FLJ23280 fis, clone HEP07194. /FEA=mRNA /DB_XREF=gi:10439907 /UG=Hs.286236 eukaryotic translation initiation factor 5 /FL=gb:AL080102.1	AL080102		NP_001960

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208708_x_at	1.26E-04	gb:AL080102.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564N1916 (from clone DKFZp564N1916); complete cds. /FEA=mRNA /GEN=DKFZp564N1916 /PROD=hypothetical protein /DB_XREF=gi:5262526 /UG=Hs.286236 eukaryotic translation initiation factor 5 /FL=gb:AL080102.1	AL080102		NP_001960
208709_s_at	5.03E-04	gb:U64898.1 /DEF=Homo sapiens NRD convertase mRNA, complete cds. /FEA=mRNA /PROD=NRD convertase /DB_XREF=gi:2897866 /UG=Hs.4099 nardilysin (N-arginine dibasic convertase) /FL=gb:U64898.1	U64898		NP_002516
208713_at	0.013223	E1B-55kDa-associated protein 5	BF724216	Hs.155218	NP_653335
208716_s_at	0.001116	gb:AB020980.1 /DEF=Homo sapiens mRNA for putative membrane protein, complete cds. /FEA=mRNA /PROD=membrane protein /DB_XREF=gi:6467174 /UG=Hs.93832 putative membrane protein /FL=gb:BC000104.1 gb:AB020980.1	AB020980		NP_061899
208723_at	0.046906	gb:BC000350.1 /DEF=Homo sapiens, ubiquitin specific protease 11, clone MGC:8620, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin specific protease 11 /DB_XREF=gi:12653164 /UG=Hs.171501 ubiquitin specific protease 11 /FL=gb:BC000350.1 gb:U44839.1 gb:NM_004651.1	BC000350		NP_004642

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208724_s_at	1.26E-04	gb:BC000905.1 /DEF=Homo sapiens, RAB1, member RAS oncogene family, clone MGC:5233, mRNA, complete cds. /FEA=mRNA /PROD=RAB1, member RAS oncogene family /DB_XREF=gi:12654174 /UG=Hs.3642 RAB1, member RAS oncogene family /FL=gb:BC000905.1	BC000905		NP_004152
208726_s_at	0.02145	gb:BC000461.1 /DEF=Homo sapiens, eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD), clone MGC:8508, mRNA, complete cds. /FEA=mRNA /PROD=eukaryotic translation initiation factor 2,subunit 2 (beta, 38kD) /DB_XREF=gi:12653382 /UG=Hs.12163 eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD) /FL=gb:BC000461.1 gb:BC000934.2 gb:M29536.1 gb:NM_003908.1	BC000461		NP_003899
208728_s_at	3.26E-04	gb:BC003682.1 /DEF=Homo sapiens, cell division cycle 42 (GTP-binding protein, 25kD), clone MGC:5044, mRNA, complete cds. /FEA=mRNA /PROD=cell division cycle 42 (GTP-binding protein,25kD) /DB_XREF=gi:13277547 /UG=Hs.146409 cell division cycle 42 (GTP-binding protein, 25kD) /FL=gb:BC002711.1 gb:BC003682.1 gb:M57298.1 gb:NM_001791.1	BC003682		NP_426359

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208729_x_at	0.02145	gb:D83043.1 /DEF=Human HLA-B mRNA, allele A*2711, complete cds. /FEA=mRNA /GEN=HLA-B /DB_XREF=gi:1871135 /UG=Hs.77961 major histocompatibility complex, class I, B /FL=gb:U36492.1 gb:U35431.1 gb:L37881.1 gb:U31971.1 gb:L42345.1 gb:U50710.1 gb:L41925.1 gb:L41214.1 gb:U27608.1 gb:U35734.1 gb:U70528.1 gb:U70529.1 gb:U80945.1 gb:U64801.1 gb:U09864.1 gb:M77776.1 gb:M77775.1 gb:L04696.1 gb:D83043.1 gb:M12678.1 gb:L09736.1 gb:L09735.1 gb:M75138.1 gb:M84381.1 gb:M84382.1 gb:M84383.1 gb:M84384.1 gb:M84385.1 gb:M32318.1 gb:M32319.1 gb:M32320.1 gb:M81798.1 gb:L04695.1 gb:M84694.1 gb:U30936.1 gb:AF016641.1 gb:U36392.1 gb:AB008102.1 gb:U75533.1 gb:U57966.1 gb:AF026218.1 gb:L07743.1 gb:U01848.1 gb:U38800.1 gb:U40498.1 gb:U03027.1 gb:U03859.1 gb:L22028.1 gb:U04243.1 gb:U06862.1 gb:L32862.1	D83043		NP_005505
208730_x_at	0.004341	Consensus includes gb:AA535244 /FEA=EST /DB_XREF=gi:2279497 /DB_XREF=est:nf93a03.s1 /CLONE=IMAGE:927436 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	NM_002865		NP_002856
208731_at	3.26E-04	Consensus includes gb:AU158062 /FEA=EST /DB_XREF=gi:11019583 /DB_XREF=est:AU158062 /CLONE=PLACE1010787 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	NM_002865		NP_002856

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208732_at	3.26E-04	Consensus includes gb:A1743756 /FEA=EST /DB_XREF=gi:5112044 /DB_XREF=est:wg53e08.x1 /CLONE=IMAGE:2368838 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	NM_002865		NP_002856
208734_x_at	0.010205	gb:M28213.1 /DEF=Homo sapiens GTP-binding protein (RAB2) mRNA, complete cds. /FEA=mRNA /GEN=RAB2 /PROD=GTP-binding protein /DB_XREF=gi:550061 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	M28213		NP_002856
208736_at	1.26E-04	gb:AF004561.1 /DEF=Homo sapiens p21-Arc mRNA, complete cds. /FEA=mRNA /PROD=p21-Arc /DB_XREF=gi:2209346 /UG=Hs.6895 actin related protein 23 complex, subunit 3 (21 kD) /FL=gb:AF004561.1 gb:AF006086.1 gb:NM_005719.1	AF004561		NP_005710
208739_x_at	3.26E-04	gb:L76416.1 /DEF=Homo sapiens MIF2 suppressor (HSMT3) mRNA, complete cds. /FEA=mRNA /GEN=HSMT3 /PROD=MIF2 suppressor /DB_XREF=gi:5566605 /UG=Hs.180139 SMT3 (suppressor of mif two 3, yeast) homolog 2 /FL=gb:L76416.1 gb:NM_006937.1	L76416		NP_008868
208741_at	0.004341	Consensus includes gb:AW274856 /FEA=EST /DB_XREF=gi:6661886 /DB_XREF=est:xm61g03.x1 /CLONE=IMAGE:2688724 /UG=Hs.23964 sin3-associated polypeptide, 18kD /FL=gb:NM_005870.2 gb:U96915.1 gb:AF153608.1 gb:U78303.1	NM_005870		NP_005861

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208742_s_at	1.26E-04	gb:U78303.1 /DEF=Human 2HOR0202 mRNA, complete cds. /FEA=mRNA /PROD=2HOR0202 /DB_XREF=gi:6648546 /UG=Hs.23964 sin3-associated polypeptide, 18kD /FL=gb:NM_005870.2 gb:U96915.1 gb:AF153608.1 gb:U78303.1	U78303		NP_005861
208743_s_at	7.58E-04	gb:BC001359.1 /DEF=Homo sapiens, tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, beta polypeptide, clone MGC:1580, mRNA, complete cds. /FEA=mRNA /PROD=tyrosine 3-monooxygenasetryptophan5-monooxygenase activation protein, beta polypeptide /DB_XREF=gi:12655022 /UG=Hs.279920 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, beta polypeptide /FL=gb:BC001359.1	BC001359		NP_647539
208746_x_at	5.03E-04	gb:AF070655.1 /DEF=Homo sapiens F1F0-type ATP synthase subunit g mRNA, complete cds. /FEA=mRNA /PROD=F1F0-type ATP synthase subunit g /DB_XREF=gi:4454685 /UG=Hs.107476 ATP synthase, H ⁺ transporting, mitochondrial F1F0, subunit g /FL=gb:AF070655.1	AF070655		NP_006467
208749_x_at	2.05E-04	gb:AF085357.1 /DEF=Homo sapiens flotillin mRNA, complete cds. /FEA=mRNA /PROD=flotillin /DB_XREF=gi:5114048 /UG=Hs.179986 flotillin 1 /FL=gb:BC001146.1 gb:AF085357.1 gb:AF089750.2 gb:NM_005803.2	AF085357		NP_005794

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208756_at	0.010205	gb:U36764.1 /DEF=Human TGF-beta receptor interacting protein 1 mRNA, complete cds. /FEA=mRNA /PROD=TGF-beta receptor interacting protein 1 /DB_XREF=gi:1036804 /UG=Hs.192023 eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD) /FL=gb:U36764.1 gb:BC000413.1 gb:BC003140.1 gb:U39067.1 gb:NM_003757.1 gb:AF116697.1	U36764		NP_003748
208757_at	0.035171	gb:BC001123.1 /DEF=Homo sapiens, Similar to gp25L2 protein, clone MGC:2142, mRNA, complete cds. /FEA=mRNA /PROD=Similar to gp25L2 protein /DB_XREF=gi:12654574 /UG=Hs.279929 gp25L2 protein /FL=gb:BC001123.1	BC001123		NP_059980
208759_at	0.003171	gb:AF240468.1 /DEF=Homo sapiens nicastrin mRNA, complete cds. /FEA=mRNA /PROD=nicastatin /DB_XREF=gi:9992877 /UG=Hs.4788 nicastrin /FL=gb:AF240468.1	AF240468		NP_056146
208761_s_at	1.26E-04	gb:U67122.1 /DEF=Human ubiquitin-related protein SUMO-1 mRNA, complete cds. /FEA=mRNA /PROD=SUMO-1 /DB_XREF=gi:1762972 /UG=Hs.81424 ubiquitin-like 1 (sentrin) /FL=gb:U61397.1 gb:U38784.1 gb:U72722.1 gb:U67122.1 gb:U83117.1 gb:NM_003352.1	U67122		NP_003343
208764_s_at	0.003171	gb:D13119.1 /DEF=Homo sapiens P2 mRNA for ATP synthase subunit c, complete cds. /FEA=mRNA /GEN=P2 /PROD=ATP synthase subunit c precursor /DB_XREF=gi:285909 /UG=Hs.89399 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 /FL=gb:D13119.1	D13119		NP_005167

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208765_s_at	0.010205	Consensus includes gb:NM_005826.1 /DEF=Homo sapiens heterogeneous nuclear ribonucleoprotein R (HNRPR), mRNA. /FEA=CDS /GEN=HNRPR /PROD=heterogeneous nuclear ribonucleoprotein R /DB_XREF=gi:5031754 /UG=Hs.15265 heterogeneous nuclear ribonucleoprotein R /FL=gb:BC001449.1 gb:AF000364.1 gb:NM_005826.1	NM_005826		NP_005817
208767_s_at	0.001116	putative integral membrane transporter	AW149681	Hs.296398	NP_060877
208770_s_at	2.05E-04	gb:BC005057.1 /DEF=Homo sapiens, eukaryotic translation initiation factor 4E binding protein 2, clone MGC:12944, mRNA, complete cds. /FEA=mRNA /PROD=eukaryotic translation initiation factor 4Ebinding protein 2 /DB_XREF=gi:13477190 /UG=Hs.278712 eukaryotic translation initiation factor 4E binding protein 2 /FL=gb:BC005057.1 gb:NM_004096.1 gb:L36056.1	BC005057		NP_004087
208771_s_at	1.26E-04	gb:J02959.1 /DEF=Human leukotriene A-4 hydrolase mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:187174 /UG=Hs.81118 leukotriene A4 hydrolase /FL=gb:J03459.1 gb:J02959.1 gb:NM_000895.1	J02959		NP_000886
208773_s_at	0.016934	gb:AL136943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586G1024 (from clone DKFZp586G1024); complete cds. /FEA=mRNA /GEN=DKFZp586G1024 /PROD=hypothetical protein /DB_XREF=gi:12053380 /UG=Hs.301226 KIAA1085 protein /FL=gb:AL136943.1	AL136943		NP_078944

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208780_x_at	1.26E-04	gb:AF154847.1 /DEF=Homo sapiens 33 kDa Vamp-associated protein (VAMP) mRNA, complete cds. /FEA=mRNA /GEN=VAMP /PROD=33 kDa Vamp-associated protein /DB_XREF=gi:8099349 /UG=Hs.9006 VAMP (vesicle-associated membrane protein)-associated protein A (33kD) /FL=gb:BC002992.1 gb:AF057358.1 gb:AF044670.1 gb:AF086627.1 gb:NM_003574.1 gb:AF154847.1	AF154847		NP_003565
208781_x_at	1.26E-04	gb:AF062483.1 /DEF=Homo sapiens SDP3 mRNA, complete cds. /FEA=mRNA /PROD=SDP3 /DB_XREF=gi:3126978 /UG=Hs.12102 sorting nexin 3 /FL=gb:AF062483.1 gb:AF034546.1 gb:NM_003795.1	AF062483		NP_690041
208783_s_at	0.016934	membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	AL570661	Hs.83532	NP_758871
208785_s_at	2.05E-04	ESTs, Highly similar to MPL3_HUMAN Microtubule-associated proteins 1A/1B light chain 3 (MAP1A/MAP1B LC3) [H.sapiens]	BE893893	Hs.356061	
208786_s_at	1.26E-04	gb:AF183417.1 /DEF=Homo sapiens microtubule-associated proteins 1A1B light chain 3 mRNA, complete cds. /FEA=mRNA /PROD=microtubule-associated proteins 1A1B lightchain 3 /DB_XREF=gi:9963772 /UG=Hs.121849 microtubule-associated proteins 1A1B light chain 3 /FL=gb:AF303888.1 gb:NM_022818.2 gb:AF183417.1	AF183417		NP_073729

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208788_at	0.007779	gb:AL136939.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1824 (from clone DKFZp586B1824); complete cds. /FEA=mRNA /GEN=DKFZp586B1824 /PROD=hypothetical protein /DB_XREF=gi:12053372 /UG=Hs.250175 homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2 /FL=gb:NM_021814.1 gb:AL136939.1 gb:AF111849.1 gb:AF231981.1	AL136939		NP_068586
208791_at	0.003026	gb:M25915.1 /DEF=Human complement cytolysis inhibitor (CLI) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:180619 /UG=Hs.75106 clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J) /FL=gb:J02908.1 gb:M25915.1 gb:M64722.1 gb:NM_001831.1	M25915		NP_001822
208792_s_at	3.26E-04	gb:M25915.1 /DEF=Human complement cytolysis inhibitor (CLI) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:180619 /UG=Hs.75106 clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J) /FL=gb:J02908.1 gb:M25915.1 gb:M64722.1 gb:NM_001831.1	M25915		NP_001822
208797_s_at	0.018206	golgin-67	AI829170	Hs.182982	NP_851422

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208799_at	1.26E-04	gb:BC004146.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, beta type, 5, clone MGC:2175, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) subunit, betatype, 5 /DB_XREF=gi:13278740 /UG=Hs.78596 proteasome (prosome, macropain) subunit, beta type, 5 /FL=gb:BC004146.1 gb:NM_002797.1 gb:D29011.1	BC004146		
208800_at	0.02145	Consensus includes gb:AV702627 /FEA=EST /DB_XREF=gi:10718957 /DB_XREF=est:AV702627 /CLONE=ADBAHC04 /UG=Hs.237825 signal recognition particle 72kD /FL=gb:AF069765.1 gb:AF077019.1 gb:NM_006947.1	NM_006947		NP_008878
208805_at	7.58E-04	gb:BC002979.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, alpha type 6, clone MGC:2333, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) subunit, alphas type 6 /DB_XREF=gi:12804240 /UG=Hs.74077 proteasome (prosome, macropain) subunit, alpha type, 6 /FL=gb:BC002979.1	BC002979		NP_002782
208807_s_at	0.016426	gb:U91543.1 /DEF=Homo sapiens zinc-finger helicase (hZFH) mRNA, complete cds. /FEA=mRNA /GEN=hZFH /PROD=zinc-finger helicase /DB_XREF=gi:3298561 /UG=Hs.25601 chromodomain helicase DNA binding protein 3 /FL=gb:U91543.1	U91543		NP_001263

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208808_s_at	1.26E-04	gb:BC000903.1 /DEF=Homo sapiens, high-mobility group (nonhistone chromosomal) protein 2, clone MGC:5234, mRNA, complete cds. /FEA=mRNA /PROD=high-mobility group (nonhistone chromosomal)protein 2 /DB_XREF=gi:12654170 /UG=Hs.80684 high-mobility group (nonhistone chromosomal) protein 2 /FL=gb:BC000903.1 gb:BC001063.1	BC000903		NP_002120
208809_s_at	0.005852	gb:AL136632.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G182 (from clone DKFZp564G182); complete cds. /FEA=mRNA /GEN=DKFZp564G182 /PROD=hypothetical protein /DB_XREF=gi:12052789 /UG=Hs.173685 hypothetical protein FLJ12619 /FL=gb:AL136632.1	AL136632		NP_112201
208810_at	0.013223	gb:AF080569.1 /DEF=Homo sapiens DnaJ-like 2 protein (HSJ2) mRNA, complete cds. /FEA=mRNA /GEN=HSJ2 /PROD=DnaJ-like 2 protein /DB_XREF=gi:4322314 /UG=Hs.181195 DnaJ (Hsp40) homolog, subfamily B, member 6 /FL=gb:AL136707.1 gb:BC000177.2 gb:AB014888.1 gb:AF080569.1 gb:NM_005494.1 gb:AF075601.1 gb:AF060703.1 gb:AB015799.1	AF080569		NP_490647
208811_s_at	0.010205	gb:AF080569.1 /DEF=Homo sapiens DnaJ-like 2 protein (HSJ2) mRNA, complete cds. /FEA=mRNA /GEN=HSJ2 /PROD=DnaJ-like 2 protein /DB_XREF=gi:4322314 /UG=Hs.181195 DnaJ (Hsp40) homolog, subfamily B, member 6 /FL=gb:AL136707.1 gb:BC000177.2 gb:AB014888.1 gb:AF080569.1 gb:NM_005494.1 gb:AF075601.1 gb:AF060703.1 gb:AB015799.1	AF080569		NP_490647

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208812_x_at	0.026891	gb:BC004489.1 /DEF=Homo sapiens, major histocompatibility complex, class I, C, clone MGC:11039, mRNA, complete cds. /FEA=mRNA /PROD=major histocompatibility complex, class I, C /DB_XREF=gi:13325360 /UG=Hs.277477 major histocompatibility complex, class I, C /FL=gb:NM_002117.1 gb:M99388.1 gb:U06487.1 gb:BC002463.1 gb:BC004489.1 gb:D64145.1 gb:D38526.1 gb:D49552.1 gb:D49819.1 gb:M24097.1 gb:M84171.1 gb:M84172.1 gb:M84173.1 gb:M84174.1 gb:M26429.1 gb:M26430.1 gb:M26431.1 gb:U41420.1 gb:U41386.1 gb:D50852.1 gb:D50853.1 gb:D50854.1 gb:D83031.1 gb:U57028.1 gb:U06695.1 gb:U06696.1 gb:M99389.1 gb:M99390.1 gb:M28160.1 gb:U09853.1 gb:AF168611.1 gb:L38251.1 gb:D31817.1	BC004489		NP_002108
208815_x_at	0.010205	gb:AB023420.1 /DEF=Homo sapiens mRNA for heat shock protein apg-2, complete cds. /FEA=mRNA /GEN=apg-2 /PROD=apg-2 /DB_XREF=gi:4579908 /UG=Hs.90093 heat shock 70kD protein 4 /FL=gb:AB023420.1	AB023420		
208818_s_at	0.02145	gb:BC000419.1 /DEF=Homo sapiens, catechol-O-methyltransferase, clone MGC:8663, mRNA, complete cds. /FEA=mRNA /PROD=catechol-O-methyltransferase /DB_XREF=gi:12653300 /UG=Hs.240013 catechol-O-methyltransferase /FL=gb:BC000419.1 gb:M58525.1 gb:M65212.1 gb:NM_007310.1 gb:NM_000754.2	BC000419		NP_009294

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208825_x_at	0.00161	gb:U43701.1 /DEF=Human ribosomal protein L23a mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L23a /DB_XREF=gi:1399085 /UG=Hs.184776 ribosomal protein L23a /FL=gb:U43701.1	U43701		NP_000975
208834_x_at	0.00161	gb:BC001865.1 /DEF=Homo sapiens, Similar to cadherin 1, type 1, E-cadherin (epithelial), clone MGC:1151, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cadherin 1, type 1, E-cadherin(epithelial) /DB_XREF=gi:12804838 /UG=Hs.194657 cadherin 1, type 1, E-cadherin (epithelial) /FL=gb:BC001865.1	BC001865		
208835_s_at	1.26E-04	cisplatin resistance-associated overexpressed protein	AW089673	Hs.3688	NP_057508
208857_s_at	1.26E-04	gb:M93008.1 /DEF=Human L-isoaspartylD-aspartyl protein carboxyl methyltransferase isozyme II mRNA, complete cds. /FEA=mRNA /PROD=L-isoaspartylD-aspartyl protein carboxylmethyltransferase /DB_XREF=gi:180636 /UG=Hs.79137 protein-L-isoaspartate (D-aspartate) O-methyltransferase /FL=gb:D25546.1 gb:M93008.1 gb:D13892.1	M93008		NP_005380
208859_s_at	0.018693	alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)	AI650257	Hs.96264	NP_612115
208864_s_at	2.05E-04	gb:AF313911.1 /DEF=Homo sapiens thioredoxin mRNA, complete cds. /FEA=mRNA /PROD=thioredoxin /DB_XREF=gi:11345419 /UG=Hs.76136 thioredoxin /FL=gb:AF313911.1 gb:BC003377.1 gb:J04026.1 gb:NM_003329.1 gb:AF276919.1 gb:AY004872.1	AF313911		NP_003320
208865_at	2.05E-04	casein kinase 1, alpha 1	BG534245	Hs.283738	NP_001883

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208867_s_at	0.033381	gb:AF119911.1 /DEF=Homo sapiens PRO2975 mRNA, complete cds. /FEA=mRNA /PROD=PRO2975 /DB_XREF=gi:7770258 /UG=Hs.144477 hypothetical protein PRO2975 /FL=gb:AF119911.1	AF119911		
208870_x_at	0.02145	gb:BC000931.2 /DEF=Homo sapiens, ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1, clone MGC:5380, mRNA, complete cds. /FEA=mRNA /PROD=ATP synthase, H+ transporting, mitochondrial F1complex, gamma polypeptide 1 /DB_XREF=gi:13111822 /UG=Hs.155433 ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1 /FL=gb:BC000470.1 gb:BC000931.2 gb:D16562.1	BC000931		NP_005165
208872_s_at	1.26E-04	likely ortholog of mouse deleted in polyposis 1	AA814140	Hs.178112	NP_005660
208873_s_at	0.005852	gb:BC000232.1 /DEF=Homo sapiens, Similar to deleted in polyposis 1, clone MGC:2267, mRNA, complete cds. /FEA=mRNA /PROD=Similar to deleted in polyposis 1 /DB_XREF=gi:12652946 /UG=Hs.178112 DNA segment, single copy probe LNS-CAILNS-CAII (deleted in polyposis /FL=gb:BC000232.1	BC000232		NP_005660
208875_s_at	0.00161	Consensus includes gb:BF796470 /FEA=EST /DB_XREF=gi:12101524 /DB_XREF=est:602259926F1 /CLONE=IMAGE:4342999 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208876_s_at	0.013223	Consensus includes gb:AI076186 /FEA=EST /DB_XREF=gi:3405364 /DB_XREF=est:oz01g01.x1 /CLONE=IMAGE:1674096 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		
208877_at	0.010205	Consensus includes gb:W74494 /FEA=EST /DB_XREF=gi:1384781 /DB_XREF=est:zd75f10.s1 /CLONE=IMAGE:346507 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		
208878_s_at	0.001116	gb:AF092132.1 /DEF=Homo sapiens PAK2 mRNA, complete cds. /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		
208880_s_at	0.009689	gb:AB019219.1 /DEF=Homo sapiens mRNA, complete cds, similar to yeast pre-mRNA splicing factors, Prp1Zer1 and Prp6. /FEA=mRNA /DB_XREF=gi:4164165 /UG=Hs.31334 putative mitochondrial outer membrane protein import receptor /FL=gb:BC001666.1 gb:AF026031.1 gb:AB019219.1 gb:NM_012469.1 gb:AF221842.1	AB019219		NP_036601
208881_x_at	1.26E-04	gb:BC005247.1 /DEF=Homo sapiens, isopentenyl-diphosphate delta isomerase, clone MGC:12281, mRNA, complete cds. /FEA=mRNA /PROD=isopentenyl-diphosphate delta isomerase /DB_XREF=gi:13528899 /UG=Hs.76038 isopentenyl-diphosphate delta isomerase /FL=gb:BC005247.1	BC005247		NP_004499
208882_s_at	3.26E-04	progesterone induced protein	U69567	Hs.278428	NP_056986

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208884_s_at	1.26E-04	gb:AF006010.1 /DEF=Homo sapiens progesterin induced protein (EDD) mRNA, complete cds. /FEA=mRNA /GEN=EDD /PROD=progesterin induced protein /DB_XREF=gi:4101694 /UG=Hs.278428 progesterin induced protein /FL=gb:NM_015902.2 gb:AF006010.1 gb:U95000.1	AF006010		NP_056986
208886_at	1.28E-04	gb:BC000145.1 /DEF=Homo sapiens, H1 histone family, member 0, clone MGC:5241, mRNA, complete cds. /FEA=mRNA /PROD=H1 histone family, member 0 /DB_XREF=gi:12652786 /UG=Hs.226117 H1 histone family, member 0 /FL=gb:BC000145.1 gb:NM_005318.1	BC000145		NP_005309
208887_at	0.005852	gb:BC000733.1 /DEF=Homo sapiens, eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD), clone MGC:2053, mRNA, complete cds. /FEA=mRNA /PROD=eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD) /DB_XREF=gi:12653882 /UG=Hs.28081 eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD) /FL=gb:AF094850.1 gb:BC000733.1 gb:AF020833.1 gb:U96074.1 gb:NM_003755.1	BC000733		NP_003746
208894_at	0.02145	gb:M60334.1 /DEF=Human MHC class II HLA-DR-alpha mRNA, complete cds. /FEA=mRNA /GEN=HLA-DRA /PROD=cell surface glycoprotein /DB_XREF=gi:188255 /UG=Hs.76807 major histocompatibility complex, class II, DR alpha /FL=gb:M60334.1 gb:NM_019111.1	M60334		NP_061984

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208895_s_at	0.00161	Consensus includes gb:BG530850 /FEA=EST /DB_XREF=gi:13522387 /DB_XREF=est:602559887F1 /CLONE=IMAGE:4698044 /UG=Hs.100555 DEADH (Asp-Glu-Ala-AspHis) box polypeptide 18 (Myc-regulated) /FL=gb:BC001238.1 gb:BC003360.1	BC003360		NP_006764
208898_at	2.05E-04	gb:AF077614.1 /DEF=Homo sapiens vacuolar ATP synthase subunit D homolog (VATD) mRNA, complete cds. /FEA=mRNA /GEN=VATD /PROD=vacuolar ATP synthase subunit D homolog /DB_XREF=gi:11999089 /UG=Hs.272630 vacuolar proton pump delta polypeptide /FL=gb:AF104629.1 gb:AF077614.1 gb:BC001411.1 gb:AF145316.1 gb:AF100741.1 gb:NM_015994.1	AF077614		NP_057078
208899_x_at	1.26E-04	gb:AF100741.1 /DEF=Homo sapiens vacuolar H-ATPase subunit D mRNA, complete cds. /FEA=mRNA /PROD=vacuolar H-ATPase subunit D /DB_XREF=gi:5138992 /UG=Hs.272630 vacuolar proton pump delta polypeptide /FL=gb:AF104629.1 gb:AF077614.1 gb:BC001411.1 gb:AF145316.1 gb:AF100741.1 gb:NM_015994.1	AF100741		NP_057078
208900_s_at	0.003539	topoisomerase (DNA) I	AW025108	Hs.317	NP_003277
208904_s_at	0.041055	gb:BC000354.1 /DEF=Homo sapiens, ribosomal protein S28, clone MGC:8658, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein S28 /DB_XREF=gi:12653172 /UG=Hs.153177 ribosomal protein S28 /FL=gb:BC000354.1	BC000354		NP_001022

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208908_s_at	0.003171	gb:AF327443.1 /DEF=Homo sapiens calpastatin mRNA, complete cds. /FEA=mRNA /PROD=calpastatin /DB_XREF=gi:12056961 /UG=Hs.279607 calpastatin /FL=gb:AF327443.1 gb:U26724.2	AF327443		NP_775085
208909_at	1.26E-04	gb:BC000649.1 /DEF=Homo sapiens, ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1, clone MGC:1362, mRNA, complete cds. /FEA=mRNA /PROD=ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1 /DB_XREF=gi:12653726 /UG=Hs.3712 ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1 /FL=gb:BC000649.1 gb:NM_006003.1	BC000649		NP_005994
208911_s_at	0.007258	gb:M34055.1 /DEF=Human pyruvate dehydrogenase E1-beta subunit mRNA, complete cds. /FEA=mRNA /GEN=PDHB /DB_XREF=gi:190791 /UG=Hs.979 pyruvate dehydrogenase (lipoamide) beta /FL=gb:BC000439.1 gb:BC001924.1 gb:J03576.1 gb:M34479.1 gb:M54788.1 gb:M34055.1 gb:NM_000925.1	M34055		NP_000916
208913_at	0.010205	Consensus includes gb:AA868560 /FEA=EST /DB_XREF=gi:2964005 /DB_XREF=est:ak43g11.s1 /CLONE=IMAGE:1408772 /UG=Hs.155546 KIAA1080 protein; Golgi-associated, gamma-adaptin ear containing, ARF-binding protein 2 /FL=gb:AF190863.1 gb:AF233522.1 gb:AF165531.1 gb:NM_015044.1	NM_015044		NP_619581

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208918_s_at	0.013223	Consensus includes gb:AI334128 /FEA=EST /DB_XREF=gi:4070687 /DB_XREF=est:qq07e12.x1 /CLONE=IMAGE:1931854 /UG=Hs.220324 hypothetical protein FLJ13052 /FL=gb:BC001709.1	BC001709		NP_075394
208919_s_at	5.03E-04	gb:BC001709.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ13052, clone MGC:1900, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ13052 /DB_XREF=gi:12804578 /UG=Hs.220324 hypothetical protein FLJ13052 /FL=gb:BC001709.1	BC001709		NP_075394
208921_s_at	2.05E-04	gb:L12387.1 /DEF=Human sorcin (SRI) mRNA, complete cds. /FEA=mRNA /GEN=SRI /PROD=sorcin /DB_XREF=gi:459835 /UG=Hs.300741 sorcin /FL=gb:M32886.1 gb:NM_003130.1 gb:L12387.1	L12387		NP_003121
208923_at	3.26E-04	gb:BC005097.1 /DEF=Homo sapiens, Similar to selective hybridizing clone, clone MGC:13167, mRNA, complete cds. /FEA=mRNA /PROD=Similar to selective hybridizing clone /DB_XREF=gi:13477256 /UG=Hs.77257 KIAA0068 protein /FL=gb:BC005097.1	BC005097		NP_055423
208924_at	3.26E-04	gb:AB024703.1 /DEF=Homo sapiens mRNA for Sid1669p, complete cds. /FEA=mRNA /GEN=Sid1669 /PROD=Sid1669p /DB_XREF=gi:5931544 /UG=Hs.96334 ring finger protein 11 /FL=gb:AF151881.1 gb:AB024703.1 gb:NM_014372.1	AB024703		NP_055187

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208926_at	8.34E-04	gb:U84246.1 /DEF=Homo sapiens lysosomal sialidase mRNA, complete cds. /FEA=mRNA /PROD=lysosomal sialidase /DB_XREF=gi:4099140 /UG=Hs.118721 sialidase 1 (lysosomal sialidase) /FL=gb:BC000722.1 gb:AF040958.1 gb:U84246.1 gb:NM_000434.1	U84246		NP_000425
208929_x_at	0.026891	gb:BC004954.1 /DEF=Homo sapiens, clone MGC:10897, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10897) /DB_XREF=gi:13436331 /UG=Hs.180842 ribosomal protein L13 /FL=gb:BC000851.1 gb:BC004954.1 gb:NM_000977.1	BC004954		NP_150254
208932_at	0.037567	gb:BC001416.1 /DEF=Homo sapiens, protein phosphatase 4 (formerly X), catalytic subunit, clone MGC:1882, mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 4 (formerly X), catalytic subunit /DB_XREF=gi:12655126 /UG=Hs.2903 protein phosphatase 4 (formerly X), catalytic subunit /FL=gb:BC001416.1 gb:AF097996.1 gb:NM_002720.1	BC001416		NP_002711
208933_s_at	0.02145	Fc fragment of IgG, low affinity IIb, receptor for (CD32)	AI659005	Hs.82316	NP_060542
208935_s_at	0.004341	gb:L78132.1 /DEF=Human prostate carcinoma tumor antigen (pcta-1) mRNA, complete cds. /FEA=mRNA /GEN=pcta-1 /PROD=prostate carcinoma tumor antigen /DB_XREF=gi:1932711 /UG=Hs.4082 lectin, galactoside-binding, soluble, 8 (galectin 8) /FL=gb:AF342815.1 gb:L78132.1 gb:AF074000.1 gb:NM_006499.1	L78132		NP_006490

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208943_s_at	0.001116	gb:U93239.1 /DEF=Human Sec62 (Sec62) mRNA, complete cds. /FEA=mRNA /GEN=Sec62 /PROD=Sec62 /DB_XREF=gi:1928972 /UG=Hs.8146 translocation protein 1 /FL=gb:D87127.1 gb:U93239.1 gb:NM_003262.1	U93239		NP_003253
208946_s_at	7.58E-04	gb:AF139131.1 /DEF=Homo sapiens beclin 1 (BECN1) mRNA, complete cds. /FEA=mRNA /GEN=BECN1 /PROD=beclin 1 /DB_XREF=gi:4680380 /UG=Hs.12272 beclin 1 (coiled-coil, myosin-like BCL2-interacting protein) /FL=gb:AF077301.1 gb:NM_003766.1 gb:AF139131.1	AF139131		NP_003757
208948_s_at	2.05E-04	gb:BC000830.1 /DEF=Homo sapiens, clone MGC:4921, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:4921) /DB_XREF=gi:12654050 /UG=Hs.6113 staufer (Drosophila, RNA-binding protein) /FL=gb:BC000830.1 gb:AF061938.1 gb:NM_017452.1	BC000830		NP_059348
208949_s_at	2.05E-04	gb:BC001120.1 /DEF=Homo sapiens, lectin, galactoside-binding, soluble, 3 (galectin 3), clone MGC:2058, mRNA, complete cds. /FEA=mRNA /PROD=lectin, galactoside-binding, soluble, 3(galectin 3) /DB_XREF=gi:12654570 /UG=Hs.621 lectin, galactoside-binding, soluble, 3 (galectin 3) /FL=gb:M35368.1 gb:BC001120.1 gb:M57710.1 gb:M36682.1 gb:AB006780.1 gb:NM_002306.1	BC001120		NP_002297

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208954_s_at	0.012367	gb:BC003381.1 /DEF=Homo sapiens, clone MGC:5353, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:5353) /DB_XREF=gi:13097242 /UG=Hs.78851 KIAA0217 protein /FL=gb:BC003381.1	BC003381		
208955_at	0.00228	gb:AB049113.1 /DEF=Homo sapiens DUT mRNA for dUTP pyrophosphatase, complete cds. /FEA=mRNA /GEN=DUT /PROD=dUTP pyrophosphatase /DB_XREF=gi:10257384 /UG=Hs.82113 dUTP pyrophosphatase /FL=gb:AB049113.1 gb:U31930.1 gb:U62891.1 gb:M89913.1 gb:NM_001948.1	AB049113		NP_001939
208959_s_at	0.004341	gb:BC005374.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 1110001E24 gene, clone MGC:12490, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 1110001E24 gene /DB_XREF=gi:13529223 /UG=Hs.154023 KIAA0573 protein /FL=gb:BC005374.1	BC005374		
208964_s_at	0.027243	gb:AL512760.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762M2311 (from clone DKFZp762M2311); complete cds. /FEA=mRNA /GEN=DKFZp762M2311 /PROD=hypothetical protein /DB_XREF=gi:12224983 /UG=Hs.132898 fatty acid desaturase 1 /FL=gb:AF084558.1 gb:NM_013402.2 gb:AL512760.1 gb:AF199596.1 gb:AF226273.1	AL512760		NP_037534
208966_x_at	0.02145	gb:AF208043.1 /DEF=Homo sapiens IFI16b (IFI16b) mRNA, complete cds. /FEA=mRNA /GEN=IFI16b /PROD=IFI16b /DB_XREF=gi:6644296 /UG=Hs.155530 interferon, gamma-inducible protein 16 /FL=gb:AF208043.1	AF208043		NP_005522

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208967_s_at	0.007779	gb:U39945.1 /DEF=Human adenylate kinase 2 (adk2) mRNA, complete cds. /FEA=mRNA /GEN=adk2 /PROD=adenylate kinase 2 /DB_XREF=gi:1209686 /UG=Hs.171811 adenylate kinase 2 /FL=gb:U39945.1 gb:U84371.1 gb:NM_001625.1	U39945		NP_751949
208970_s_at	7.58E-04	gb:M14016.1 /DEF=Human uroporphyrinogen decarboxylase mRNA, complete cds. /FEA=mRNA /GEN=UROD /DB_XREF=gi:340180 /UG=Hs.78601 uroporphyrinogen decarboxylase /FL=gb:BC001778.1 gb:M14016.1 gb:AF104421.1 gb:AF104422.1 gb:AF104423.1 gb:AF104424.1 gb:AF104425.1 gb:AF104426.1 gb:AF104427.1 gb:AF104428.1 gb:AF104429.1 gb:AF104430.1 gb:AF104431.1 gb:AF104432.1 gb:AF104433.1 gb:AF104434.1 gb:AF104435.1 gb:AF104436.1 gb:AF104437.1 gb:AF104438.1 gb:AF104439.1 gb:AF104440.1 gb:NM_000374.2	M14016		NP_000365
208977_x_at	0.00161	gb:BC004188.1 /DEF=Homo sapiens, tubulin, beta, 2, clone MGC:2826, mRNA, complete cds. /FEA=mRNA /PROD=tubulin, beta, 2 /DB_XREF=gi:13278848 /UG=Hs.251653 tubulin, beta, 2 /FL=gb:BC002783.1 gb:BC002885.1 gb:BC001911.1 gb:BC004188.1 gb:NM_006088.1	BC004188		NP_006079

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208979_at	1.26E-04	gb:AF128458.1 /DEF=Homo sapiens nuclear receptor coactivator RAP250 mRNA, complete cds. /FEA=mRNA /PROD=nuclear receptor coactivator RAP250 /DB_XREF=gi:7140573 /UG=Hs.159613 thyroid hormone receptor binding protein /FL=gb:AF177388.1 gb:AF208227.1 gb:AF128458.1 gb:NM_014071.1 gb:AF171667.1 gb:AF245115.1	AF128458		NP_054790
208981_at	1.26E-04	platelet/endothelial cell adhesion molecule (CD31 antigen)	AW574504	Hs.78146	NP_000433
208982_at	7.58E-04	platelet/endothelial cell adhesion molecule (CD31 antigen)	AW574504	Hs.78146	NP_000433
208988_at	0.033381	Consensus includes gb:BE675843 /FEA=EST /DB_XREF=gi:10036384 /DB_XREF=est:7f17b04.x1 /CLONE=IMAGE:3294895 /UG=Hs.219614 f-box and leucine rich repeat protein 11 /FL=gb:AF179221.1	AK024505		NP_036440
208989_s_at	0.003529	gb:AF179221.1 /DEF=Homo sapiens F-box protein Lilina (LILINA) mRNA, complete cds. /FEA=mRNA /GEN=LILINA /PROD=F-box protein Lilina /DB_XREF=gi:5917729 /UG=Hs.219614 f-box and leucine rich repeat protein 11 /FL=gb:AF179221.1	AF179221		NP_036440
208991_at	0.010205	signal transducer and activator of transcription 3 (acute-phase response factor)	AA634272	Hs.321677	
208993_s_at	6.68E-04	Consensus includes gb:AW340788 /FEA=EST /DB_XREF=gi:6837414 /DB_XREF=est:ha12h05.x1 /CLONE=IMAGE:2873529 /UG=Hs.77965 peptidyl-prolyl isomerase G (cyclophilin G) /FL=gb:U40763.1 gb:NM_004792.1	NM_004792		NP_004783

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208994_s_at	7.58E-04	Consensus includes gb:AI638762 /FEA=EST /DB_XREF=gi:4690996 /DB_XREF=est:tt32a04.x1 /CLONE=IMAGE:2242446 /UG=Hs.77965 peptidyl-prolyl isomerase G (cyclophilin G) /FL=gb:U40763.1 gb:NM_004792.1	NM_004792		NP_004783
209003_at	0.007779	gb:AF070548.1 /DEF=Homo sapiens clone 24408 2-oxoglutarate carrier protein mRNA, complete cds. /FEA=mRNA /PROD=2-oxoglutarate carrier protein /DB_XREF=gi:3387910 /UG=Hs.184877 solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11 /FL=gb:AF070548.1	AF070548		NP_003553
209004_s_at	0.005852	gb:AF142481.1 /DEF=Homo sapiens F-box protein FLR1 (FLR1) mRNA, complete cds. /FEA=mRNA /GEN=FLR1 /PROD=F-box protein FLR1 /DB_XREF=gi:7672733 /UG=Hs.5548 f-box and leucine-rich repeat protein 5 /FL=gb:AF199420.1 gb:AF142481.1 gb:AF157323.1	AF142481		NP_277077
209005_at	0.00228	gb:AF157323.1 /DEF=Homo sapiens p45SKP2-like protein mRNA, complete cds. /FEA=mRNA /PROD=p45SKP2-like protein /DB_XREF=gi:7688696 /UG=Hs.5548 f-box and leucine-rich repeat protein 5 /FL=gb:AF199420.1 gb:AF142481.1 gb:AF157323.1	AF157323		NP_277077

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209015_s_at	1.26E-04	gb:BC002446.1 /DEF=Homo sapiens, MRJ gene for a member of the DNAJ protein family, clone MGC:1152, mRNA, complete cds. /FEA=mRNA /PROD=MRJ gene for a member of the DNAJ proteinfamily /DB_XREF=gi:12803262 /UG=Hs.181195 DnaJ (Hsp40) homolog, subfamily B, member 6 /FL=gb:BC002446.1 gb:AB015798.1	BC002446		NP_490647
209018_s_at	7.58E-04	PTEN induced putative kinase 1	BF432478	Hs.6163	NP_115785
209020_at	5.03E-04	gb:AF217514.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM038 mRNA, complete cds. /FEA=mRNA /PROD=uncharacterized bone marrow protein BM038 /DB_XREF=gi:7688970 /UG=Hs.75798 hypothetical protein /FL=gb:AF161517.1 gb:AF151041.1 gb:AF217514.1 gb:NM_016470.1	AF217514		NP_057554
209022_at	0.003171	Consensus includes gb:AK026678.1 /DEF=Homo sapiens cDNA: FLJ23025 fis, clone LNG01702, highly similar to HUMAUTOND Human autonomously replicating sequence (ARS) mRNA. /FEA=mRNA /DB_XREF=gi:10439584 /UG=Hs.8217 stromal antigen 2 /FL=gb:BC001765.1	AK026678		NP_006594
209024_s_at	0.002418	NS1-associated protein 1	AI472757	Hs.373499	NP_006363
209025_s_at	0.003171	gb:AF037448.1 /DEF=Homo sapiens RRM RNA binding protein Gry-rbp (GRY-RBP) mRNA, complete cds. /FEA=mRNA /GEN=GRY-RBP /PROD=Gry-rbp /DB_XREF=gi:3037012 /UG=Hs.155489 NS1-associated protein 1 /FL=gb:AF037448.1	AF037448		NP_006363
209027_s_at	3.26E-04	spectrin SH3 domain binding protein 1	BF673013	Hs.24752	NP_005461

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209028_s_at	0.026891	gb:AF006516.1 /DEF=Homo sapiens eps8 binding protein e3B1 mRNA, complete cds. /FEA=mRNA /PROD=e3B1 /DB_XREF=gi:2245670 /UG=Hs.24752 spectrin SH3 domain binding protein 1 /FL=gb:AF006516.1	AF006516		NP_005461
209029_at	0.003171	gb:AF193844.1 /DEF=Homo sapiens COP9 complex subunit 7a mRNA, complete cds. /FEA=mRNA /PROD=COP9 complex subunit 7a /DB_XREF=gi:6120134 /UG=Hs.3758 COP9 complex subunit 7a /FL=gb:AB033603.1 gb:AF193844.1 gb:AF210052.1 gb:NM_016319.1	AF193844		NP_057403
209033_s_at	0.001116	gb:D86550.1 /DEF=Human mRNA for serinethreonine protein kinase, complete cds. /FEA=mRNA /GEN=hMNB /PROD=serinethreonine protein kinase /DB_XREF=gi:1772437 /UG=Hs.75842 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A /FL=gb:D85759.1 gb:D86550.1	D86550		NP_569122
209034_at	7.58E-04	gb:AF279899.1 /DEF=Homo sapiens PNAS-145 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-145 /DB_XREF=gi:12751123 /UG=Hs.75969 proline-rich protein with nuclear targeting signal /FL=gb:AF279899.1 gb:U03105.1 gb:NM_006813.1	AF279899		NP_006804

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209043_at	0.004341	gb:AF033026.1 /DEF=Homo sapiens bifunctional ATP sulfurylaseadenosine 5-phosphosulfate kinase mRNA, complete cds. /FEA=mRNA /PROD=bifunctional ATP sulfurylaseadenosine5-phosphosulfate kinase /DB_XREF=gi:3378100 /UG=Hs.3833 3-phosphoadenosine 5-phosphosulfate synthase 1 /FL=gb:AF033026.1 gb:AF016496.1 gb:NM_005443.1 gb:AF105227.1	AF033026		NP_005434
209044_x_at	0.003171	gb:BC004273.1 /DEF=Homo sapiens, splicing factor 3b, subunit 4, 49kD, clone MGC:10828, mRNA, complete cds. /FEA=mRNA /PROD=splicing factor 3b, subunit 4, 49kD /DB_XREF=gi:13279088 /UG=Hs.25797 splicing factor 3b, subunit 4, 49kD /FL=gb:BC004273.1 gb:NM_005850.1	BC004273		NP_005841
209046_s_at	1.26E-04	gb:AB030710.1 /DEF=Homo sapiens FLC3A mRNA for MAP1 light chain 3 related protein, complete cds. /FEA=mRNA /GEN=FLC3A /PROD=MAP1 light chain 3 related protein /DB_XREF=gi:12641848 /UG=Hs.6518 ganglioside expression factor 2 /FL=gb:AB030710.1 gb:AF087848.1 gb:AF077046.1 gb:NM_007285.1	AB030710		NP_009216
209050_s_at	0.026891	ral guanine nucleotide dissociation stimulator	AI421559	Hs.396157	NP_006257

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209054_s_at	0.004341	gb:AF083389.1 /DEF=Homo sapiens putative WHSC1 protein (WHSC1) mRNA, alternative splice product with complete exon 12, complete cds. /FEA=mRNA /GEN=WHSC1 /PROD=putative WHSC1 protein /DB_XREF=gi:4378016 /UG=Hs.110457 Wolf-Hirschhorn syndrome candidate 1 /FL=gb:AF083389.1	AF083389		NP_579891
209056_s_at	0.016934	CDC5 cell division cycle 5-like (S. pombe)	AW268817	Hs.155174	NP_001244
209059_s_at	0.033381	gb:AB002282.1 /DEF=Homo sapiens mRNA for hMBF1alpha, complete cds. /FEA=mRNA /PROD=hMBF1alpha /DB_XREF=gi:6526354 /UG=Hs.174050 endothelial differentiation-related factor 1 /FL=gb:NM_003792.1 gb:AB002282.1	AB002282		NP_694880
209060_x_at	0.001054	Consensus includes gb:AI438999 /FEA=EST /DB_XREF=gi:4301251 /DB_XREF=est:tc84b12.x1 /CLONE=IMAGE:2072831 /UG=Hs.225977 nuclear receptor coactivator 3 /FL=gb:AF010227.1 gb:AF012108.1 gb:AF016031.1	AF012108		NP_006525
209061_at	0.013223	Consensus includes gb:AI761748 /FEA=EST /DB_XREF=gi:5177504 /DB_XREF=est:wg67h01.x1 /CLONE=IMAGE:2370193 /UG=Hs.225977 nuclear receptor coactivator 3 /FL=gb:AF010227.1 gb:AF012108.1 gb:AF016031.1	AF012108		NP_006525
209063_x_at	0.00228	ESTs, Highly similar to polyadenylate binding protein-interacting protein 1 [Homo sapiens] [H.sapiens]	BF248165	Hs.374614	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209067_s_at	0.016934	gb:D89092.1 /DEF=Homo sapiens hnRNP JKTBP mRNA, complete cds. /FEA=mRNA /GEN=hnRNP JKTBP /DB_XREF=gi:2780747 /UG=Hs.170311 heterogeneous nuclear ribonucleoprotein D-like /FL=gb:D89092.1 gb:D89678.1	D89092		NP_112740
209073_s_at	3.26E-04	gb:AF015040.1 /DEF=Homo sapiens NUMB protein (NUMB) mRNA, complete cds. /FEA=mRNA /GEN=NUMB /PROD=NUMB protein /DB_XREF=gi:4102704 /UG=Hs.78890 numb (Drosophila) homolog /FL=gb:AF015040.1 gb:AF108092.1 gb:AF171939.1	AF015040		NP_003735
209076_s_at	0.001116	gb:BC000974.2 /DEF=Homo sapiens, Similar to hypothetical protein 628, clone MGC:5116, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein 628 /DB_XREF=gi:12803025 /UG=Hs.181349 hypothetical protein 628 /FL=gb:BC000974.2	BC000974		NP_062559
209080_x_at	0.02145	gb:AF118652.1 /DEF=Homo sapiens PKCq-interacting protein PICOT (PICOT) mRNA, complete cds. /FEA=mRNA /GEN=PICOT /PROD=PKCq-interacting protein PICOT /DB_XREF=gi:6840952 /UG=Hs.42644 thioredoxin-like /FL=gb:BC005289.1 gb:AF118649.1 gb:AF118652.1	AF118652		NP_006532
209088_s_at	6.68E-04	ubinnuclein 1	T70262	Hs.21479	NP_058632
209089_at	7.58E-04	gb:BC001267.1 /DEF=Homo sapiens, RAB5A, member RAS oncogene family, clone MGC:5048, mRNA, complete cds. /FEA=mRNA /PROD=RAB5A, member RAS oncogene family /DB_XREF=gi:12654846 /UG=Hs.73957 RAB5A, member RAS oncogene family /FL=gb:BC001267.1	BC001267		NP_004153

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209090_s_at	2.05E-04	Consensus includes gb:AL049597 /DEF=Human DNA sequence from clone RP4-612B15 on chromosome 1p22.2-31.1. Contains the (possibly pseudo) gene for a novel protein similar to 60S ribosomal protein L17 (RPL17), the gene for CGI-61, endophilin B1 and KIAA0491, ESTs, STSs, GSSs and two CpG... /FEA=mRNA_3 /DB_XREF=gi:9663352 /UG=Hs.136309 SH3-containing protein SH3GLB1 /FL=gb:AF263293.1	AL049597		
209091_s_at	1.26E-04	gb:AF263293.1 /DEF=Homo sapiens endophilin B1 mRNA, complete cds. /FEA=mRNA /PROD=endophilin B1 /DB_XREF=gi:8118529 /UG=Hs.136309 SH3-containing protein SH3GLB1 /FL=gb:AF263293.1	AF263293		NP_057093
209093_s_at	0.026891	gb:K02920.1 /DEF=Human lysosomal glucocerebrosidase mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:183021 /UG=Hs.282997 glucosidase, beta; acid (includes glucosylceramidase) /FL=gb:BC003356.1 gb:M16328.1 gb:K02920.1 gb:M19285.1 gb:NM_000157.1 gb:D13286.1	K02920		NP_000148
209096_at	0.02145	gb:U62136.2 /DEF=Homo sapiens enterocyte differentiation associated factor EDAF-1 mRNA, complete cds. /FEA=mRNA /PROD=enterocyte differentiation associated factor EDAF-1 /DB_XREF=gi:4775663 /UG=Hs.79300 ubiquitin-conjugating enzyme E2 variant 2 /FL=gb:NM_003350.2 gb:AF049140.1 gb:U62136.2	U62136		NP_003341

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209098_s_at	0.041795	gb:U61276.1 /DEF=Human transmembrane protein Jagged 1 (HJ1) mRNA, complete cds. /FEA=mRNA /GEN=HJ1 /PROD=transmembrane protein Jagged 1 /DB_XREF=gi:1438936 /UG=Hs.91143 jagged 1 (Alagille syndrome) /FL=gb:U61276.1 gb:U73936.1 gb:AF003837.1 gb:AF028593.1 gb:NM_000214.1	U61276		NP_000205
209099_x_at	1.28E-04	gb:U73936.1 /DEF=Homo sapiens Jagged 1 (HJ1) mRNA, complete cds. /FEA=mRNA /GEN=HJ1 /PROD=Jagged 1 /DB_XREF=gi:1695273 /UG=Hs.91143 jagged 1 (Alagille syndrome) /FL=gb:U61276.1 gb:U73936.1 gb:AF003837.1 gb:AF028593.1 gb:NM_000214.1	U73936		NP_000205
209102_s_at	1.26E-04	gb:AF019214.1 /DEF=Homo sapiens HMG box containing protein 1 mRNA, complete cds. /FEA=mRNA /PROD=HMG box containing protein 1 /DB_XREF=gi:2460168 /UG=Hs.10882 HMG-box containing protein 1 /FL=gb:AF019214.1	AF019214		NP_036389
209103_s_at	0.001116	gb:BC001049.1 /DEF=Homo sapiens, Similar to ubiquitin fusion degradation 1 like, clone MGC:1385, mRNA, complete cds. /FEA=mRNA /PROD=Similar to ubiquitin fusion degradation 1 like /DB_XREF=gi:12654446 /UG=Hs.181369 ubiquitin fusion degradation 1-like /FL=gb:BC001049.1 gb:BC005087.1 gb:AF141201.1	BC001049		NP_005650
209105_at	0.026891	Consensus includes gb:AI672428 /FEA=EST /DB_XREF=gi:4852159 /DB_XREF=est:wa03a11.x1 /CLONE=IMAGE:2296988 /UG=Hs.74002 nuclear receptor coactivator 1 /FL=gb:U19179.1	U19179		NP_671766

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209111_at	0.044231	gb:BC004155.1 /DEF=Homo sapiens, ring finger protein 5, clone MGC:2407, mRNA, complete cds. /FEA=mRNA /PROD=ring finger protein 5 /DB_XREF=gi:13278761 /UG=Hs.216354 ring finger protein 5 /FL=gb:BC004155.1 gb:AB056869.1 gb:NM_006913.1	BC004155		NP_008844
209112_at	0.010205	gb:BC001971.1 /DEF=Homo sapiens, Similar to cyclin-dependent kinase inhibitor 1B (p27, Kip1), clone MGC:5304, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cyclin-dependent kinase inhibitor 1B(p27, Kip1) /DB_XREF=gi:12805034 /UG=Hs.238990 cyclin-dependent kinase inhibitor 1B (p27, Kip1) /FL=gb:BC001971.1 gb:NM_004064.1 gb:U10906.1 gb:AF247551.1 gb:AY004255.1	BC001971		NP_004055
209115_at	0.010205	gb:AL117566.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566J164 (from clone DKFZp566J164); complete cds. /FEA=mRNA /GEN=DKFZp566J164 /PROD=hypothetical protein /DB_XREF=gi:5912116 /UG=Hs.154320 ubiquitin-activating enzyme E1C (homologous to yeast UBA3) /FL=gb:AF046024.1 gb:AB012190.1 gb:NM_003968.1 gb:AL117566.1	AL117566		NP_003959
209116_x_at	3.56E-04	gb:M25079.1 /DEF=Human sickle cell beta-globin mRNA, complete cds. /FEA=mRNA /PROD=beta-globin /DB_XREF=gi:179408 /UG=Hs.155376 hemoglobin, beta /FL=gb:NM_000518.3 gb:M25079.1 gb:AF117710.1 gb:AF181989.1	M25079		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209117_at	7.58E-04	gb:U79458.1 /DEF=Human WW domain binding protein-2 mRNA, complete cds. /FEA=mRNA /PROD=WW domain binding protein-2 /DB_XREF=gi:4205085 /UG=Hs.231840 WW domain binding protein 2 /FL=gb:U79458.1	U79458		
209124_at	2.05E-04	gb:U70451.1 /DEF=Human myeloid differentiation primary response protein MyD88 mRNA, complete cds. /FEA=mRNA /PROD=MyD88 /DB_XREF=gi:1763090 /UG=Hs.82116 myeloid differentiation primary response gene (88) /FL=gb:U70451.1 gb:U84408.1 gb:NM_002468.1	U70451		NP_002459
209127_s_at	0.035763	Consensus includes gb:AW173076 /FEA=EST /DB_XREF=gi:6439024 . /DB_XREF=est:xj82h07.x1 /CLONE=IMAGE:2663773 /UG=Hs.116875 KIAA0156 gene product /FL=gb:AB020880.1 gb:NM_014706.1 gb:D63879.1	NM_014706		NP_055521
209129_at	0.00423	gb:AF000974.1 /DEF=Human zyxin related protein ZRP-1 mRNA, complete cds. /FEA=mRNA /PROD=ZRP-1 /DB_XREF=gi:2232135 /UG=Hs.119498 thyroid hormone receptor interactor 6 /FL=gb:BC002680.1 gb:BC004249.1 gb:BC004999.1 gb:AF000974.1	AF000974		NP_003293
209130_at	0.00161	gb:BC003686.1 /DEF=Homo sapiens, synaptosomal-associated protein, 23kD, clone MGC:5155, mRNA, complete cds. /FEA=mRNA /PROD=synaptosomal-associated protein, 23kD /DB_XREF=gi:13277555 /UG=Hs.184376 synaptosomal-associated protein, 23kD /FL=gb:BC000148.2 gb:BC003686.1 gb:U55936.1 gb:Y09567.1	BC003686		NP_570710

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209140_x_at	0.004341	gb:L42024.1 /DEF=Homo sapiens MHC HLA-B39 mRNA, complete cds. /FEA=mRNA /GEN=HLA-B39 /PROD=major histocompatibility complex /DB_XREF=gi:804748 /UG=Hs.77961 major histocompatibility complex, class I, B /FL=gb:L37880.1 gb:U29057.1 gb:U29480.1 gb:U63653.1 gb:D85761.1 gb:D85762.1 gb:M77774.1 gb:M77778.1 gb:M77777.1 gb:M16102.1 gb:M84380.1 gb:M32317.1 gb:M24033.1 gb:L17005.1 gb:U88407.1 gb:U29083.1 gb:L24373.1 gb:U03698.1 gb:U04787.1 gb:U04244.1 gb:U04245.1 gb:NM_005514.1 gb:L33922.1 gb:AF189017.1 gb:U21052.1 gb:U21053.1 gb:L36318.1 gb:U09912.1 gb:L42024.1	L42024		NP_005505
209141_at	2.05E-04	ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)	AW299555	Hs.78563	NP_003333
209142_s_at	1.26E-04	gb:BC002775.1 /DEF=Homo sapiens, Similar to CG9602 gene product, clone MGC:3733, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG9602 gene product /DB_XREF=gi:12803862 /UG=Hs.78563 ubiquitin-conjugating enzyme E2G 1 (homologous to C. elegans UBC7) /FL=gb:BC002775.1 gb:NM_003342.1 gb:D78514.1	BC002775		NP_003333
209146_at	0.016934	sterol-C4-methyl oxidase-like	AV704962	Hs.239926	NP_006736
209149_s_at	0.029453	transmembrane 9 superfamily member 1	BE899402	Hs.91586	NP_006396
209150_s_at	1.26E-04	gb:U94831.1 /DEF=Homo sapiens multispanning membrane protein mRNA, complete cds. /FEA=mRNA /PROD=multispanning membrane protein /DB_XREF=gi:2276459 /UG=Hs.91586 transmembrane 9 superfamily member 1 /FL=gb:U94831.1 gb:NM_006405.1	U94831		NP_006396

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209154_at	0.007779	gb:AF234997.1 /DEF=Homo sapiens glutaminase-interacting protein 3 mRNA, complete cds. /FEA=mRNA /PROD=glutaminase-interacting protein 3 /DB_XREF=gi:12005281 /UG=Hs.12956 Tax interaction protein 1 /FL=gb:AF028823.2 gb:NM_014604.1 gb:AF234997.1	AF234997		NP_055419
209155_s_at	1.26E-04	gb:BC001595.1 /DEF=Homo sapiens, 5-nucleotidase (purine), cytosolic type B, clone MGC:1109, mRNA, complete cds. /FEA=mRNA /PROD=5-nucleotidase (purine), cytosolic type B /DB_XREF=gi:12804388 /UG=Hs.138593 5-nucleotidase (purine), cytosolic type B /FL=gb:BC001595.1	BC001595		NP_036361
209165_at	0.010205	gb:AF083208.1 /DEF=Homo sapiens Che-1 mRNA, complete cds. /FEA=mRNA /PROD=Che-1 /DB_XREF=gi:5813798 /UG=Hs.16178 apoptosis antagonizing transcription factor /FL=gb:BC000591.1 gb:AF083208.1 gb:NM_012138.1	AF083208		NP_036270
209166_s_at	0.013223	gb:U68567.1 /DEF=Human lysosomal acid alpha-mannosidase mRNA, complete cds. /FEA=mRNA /PROD=lysosomal acid alpha-mannosidase /DB_XREF=gi:1658373 /UG=Hs.279854 mannosidase, alpha, class 2B, member 1 /FL=gb:NM_000528.1 gb:BC000736.1 gb:U05572.1 gb:U68567.1 gb:U60266.1	U68567		NP_000519

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209175_at	0.013223	Consensus includes gb:AK001135.1 /DEF=Homo sapiens cDNA FLJ10273 fis, clone HEMBB1001137, highly similar to Homo sapiens mRNA for putative phospholipase. /FEA=mRNA /DB_XREF=gi:7022202 /UG=Hs.300208 Sec23-interacting protein p125 /FL=gb:BC002540.1 gb:AB019435.1 gb:NM_007190.1	AK001135		NP_009121
209185_s_at	7.58E-04	gb:AF073310.1 /DEF=Homo sapiens insulin receptor substrate-2 (IRS2) mRNA, complete cds. /FEA=mRNA /GEN=IRS2 /PROD=insulin receptor substrate-2 /DB_XREF=gi:4511968 /UG=Hs.143648 insulin receptor substrate 2 /FL=gb:NM_003749.1 gb:AF073310.1	AF073310		NP_003740
209186_at	0.00161	gb:M23114.1 /DEF=Homo sapiens calcium-ATPase (HK1) mRNA, complete cds. /FEA=mRNA /GEN=HK1 /DB_XREF=gi:184100 /UG=Hs.1526 ATPase, Ca++ transporting, cardiac muscle, slow twitch 2 /FL=gb:M23114.1	M23114		NP_733765
209187_at	7.58E-04	down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AW516932	Hs.283643	NP_001929
209188_x_at	1.26E-04	gb:BC002809.1 /DEF=Homo sapiens, down-regulator of transcription 1, TBP-binding (negative cofactor 2), clone MGC:4292, mRNA, complete cds. /FEA=mRNA /PROD=down-regulator of transcription 1, TBP-binding(negative cofactor 2) /DB_XREF=gi:12803924 /UG=Hs.16697 down-regulator of transcription 1, TBP-binding (negative cofactor 2) /FL=gb:BC002809.1	BC002809		NP_001929

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209190_s_at	5.03E-04	gb:AF051782.1 /DEF=Homo sapiens diaphanous 1 (HDIA1) mRNA, complete cds. /FEA=mRNA /GEN=HDIA1 /PROD=diaphanous 1 /DB_XREF=gi:2947237 /UG=Hs.26584 diaphanous (Drosophila, homolog) 1 /FL=gb:AF051782.1 gb:NM_005219.1	AF051782		NP_005210
209194_at	3.26E-04	gb:BC005334.1 /DEF=Homo sapiens, centrin, EF-hand protein, 2, clone MGC:12421, mRNA, complete cds. /FEA=mRNA /PROD=centrin, EF-hand protein, 2 /DB_XREF=gi:13529121 /UG=Hs.82794 centrin, EF-hand protein, 2 /FL=gb:BC005334.1 gb:NM_004344.1	BC005334		NP_004335
209198_s_at	0.033381	gb:BC004291.1 /DEF=Homo sapiens, Similar to synaptotagmin 11, clone MGC:10881, mRNA, complete cds. /FEA=mRNA /PROD=Similar to synaptotagmin 11 /DB_XREF=gi:13279139 /UG=Hs.74554 KIAA0080 protein /FL=gb:BC004291.1	BC004291		NP_689493
209199_s_at	0.007779	MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	N22468	Hs.78995	NP_002388
209202_s_at	0.002536	gb:AF001690.1 /DEF=Homo sapiens EXT like protein 3 (EXTL3) mRNA, complete cds. /FEA=mRNA /GEN=EXTL3 /PROD=EXT like protein 3 /DB_XREF=gi:2897904 /UG=Hs.9018 exostoses (multiple) like 3 /FL=gb:AB007042.1 gb:AF001690.1 gb:AB011091.1 gb:AF029231.1 gb:NM_001440.1 gb:AF083551.1	AF001690		NP_001431
209206_at	2.05E-04	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	AV701283	Hs.50785	NP_004883

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209211_at	1.49E-04	gb:AF132818.1 /DEF=Homo sapiens colon Kruppel-like factor (CKLF) mRNA, complete cds. /FEA=mRNA /GEN=CKLF /PROD=colon Kruppel-like factor /DB_XREF=gi:6580834 /UG=Hs.84728 Kruppel-like factor 5 (intestinal) /FL=gb:D14520.1 gb:NM_001730.1 gb:AF132818.1 gb:AB030824.1 gb:AF287272.1	AF132818		NP_001721
209214_s_at	0.02145	gb:BC004817.1 /DEF=Homo sapiens, Ewing sarcoma breakpoint region 1, clone MGC:5432, mRNA, complete cds. /FEA=mRNA /PROD=Ewing sarcoma breakpoint region 1 /DB_XREF=gi:13435962 /UG=Hs.129953 Ewing sarcoma breakpoint region 1 /FL=gb:BC004817.1 gb:NM_005243.1	BC004817		NP_053733
209215_at	0.005039	gb:L11669.1 /DEF=Human tetracycline transporter-like protein mRNA, complete cds. /FEA=mRNA /PROD=tetracycline transporter-like protein /DB_XREF=gi:307501 /UG=Hs.157145 tetracycline transporter-like protein /FL=gb:L11669.1 gb:NM_001120.1	L11669		NP_001111
209216_at	2.05E-04	gb:BC000464.1 /DEF=Homo sapiens, JM5 protein, clone MGC:8542, mRNA, complete cds. /FEA=mRNA /PROD=JM5 protein /DB_XREF=gi:12653388 /UG=Hs.21753 JM5 protein /FL=gb:BC000464.1 gb:BC003037.1 gb:AJ005897.1 gb:NM_007075.1	BC000464		NP_009006

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209217_s_at	1.26E-04	gb:BC000464.1 /DEF=Homo sapiens, JM5 protein, clone MGC:8542, mRNA, complete cds. /FEA=mRNA /PROD=JM5 protein /DB_XREF=gi:12653388 /UG=Hs.21753 JM5 protein /FL=gb:BC000464.1 gb:BC003037.1 gb:AJ005897.1 gb:NM_007075.1	BC000464		NP_009006
209218_at	0.00228	gb:AF098865.1 /DEF=Homo sapiens squalene epoxidase (ERG1) mRNA, complete cds. /FEA=mRNA /GEN=ERG1 /PROD=squalene epoxidase /DB_XREF=gi:4204674 /UG=Hs.71465 squalene epoxidase /FL=gb:D78130.1 gb:AF098865.1 gb:NM_003129.2	AF098865		NP_003120
209219_at	0.021312	gb:L03411.1 /DEF=Human RD protein (RD) mRNA, complete cds. /FEA=mRNA /GEN=RD /PROD=RD protein /DB_XREF=gi:190973 /UG=Hs.106061 RD RNA-binding protein /FL=gb:L03411.1 gb:NM_002904.1	L03411		NP_002895
209221_s_at	0.001116	oxysterol binding protein-like 2	AI753638	Hs.15519	NP_653081
209224_s_at	1.26E-04	gb:BC003674.1 /DEF=Homo sapiens, NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8), clone MGC:12315, mRNA, complete cds. /FEA=mRNA /PROD=NADH dehydrogenase (ubiquinone) 1 alphasubcomplex, 2 (8kD, B8) /DB_XREF=gi:13277539 /UG=Hs.163867 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8) /FL=gb:BC003674.1 gb:AF047185.1 gb:NM_002488.1 gb:AF077029.1	BC003674		NP_002479
209234_at	1.26E-04	kinesin family member 1B	BF939474	Hs.375193	NP_055889
209238_at	2.05E-04	syntaxin 3A	BE966922	Hs.82240	NP_004168

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209239_at	5.03E-04	gb:M55643.1 /DEF=Human factor KBF1 mRNA, complete cds. /FEA=mRNA /GEN=NF-kappa-B /PROD=factor KBF1 /DB_XREF=gi:189179 /UG=Hs.83428 nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105) /FL=gb:NM_003998.1 gb:M58603.1 gb:M55643.1	M55643		NP_003989
209247_s_at	0.028552	gb:BC001661.1 /DEF=Homo sapiens, ATP-binding cassette, sub-family F (GCN20), member 2, clone MGC:1425, mRNA, complete cds. /FEA=mRNA /PROD=ATP-binding cassette, sub-family F (GCN20), member 2 /DB_XREF=gi:12804500 /UG=Hs.153612 ATP-binding cassette, sub-family F (GCN20), member 2 /FL=gb:AF261091.1 gb:AF261092.1 gb:BC001661.1 gb:AF091073.1	BC001661		NP_009120
209248_at	0.00228	gb:AL136713.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566K113 (from clone DKFZp566K113); complete cds. /FEA=mRNA /GEN=DKFZp566K113 /PROD=hypothetical protein /DB_XREF=gi:12052945 /UG=Hs.182470 PTD010 protein /FL=gb:AF060923.1 gb:AL136713.1 gb:AF131820.1 gb:AF078863.1 gb:AB009685.1 gb:NM_014394.1	AL136713		NP_055209
209249_s_at	2.05E-04	gb:AF131820.1 /DEF=Homo sapiens clone 25077 mRNA sequence, complete cds. /FEA=mRNA /PROD=Unknown /DB_XREF=gi:4406655 /UG=Hs.182470 PTD010 protein /FL=gb:AF060923.1 gb:AL136713.1 gb:AF131820.1 gb:AF078863.1 gb:AB009685.1 gb:NM_014394.1	AF131820		NP_055209

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209250_at	0.001116	gb:BC000961.2 /DEF=Homo sapiens, degenerative spermatocyte (homolog Drosophila; lipid desaturase), clone MGC:5079, mRNA, complete cds. /FEA=mRNA /PROD=degenerative spermatocyte (homolog Drosophila; lipid desaturase) /DB_XREF=gi:12803018 /UG=Hs.185973 degenerative spermatocyte (homolog Drosophila; lipid desaturase) /FL=gb:BC000961.2	BC000961		NP_659004
209251_x_at	3.26E-04	gb:BC004949.1 /DEF=Homo sapiens, tubulin alpha 6, clone MGC:10851, mRNA, complete cds. /FEA=mRNA /PROD=tubulin alpha 6 /DB_XREF=gi:13436316 /UG=Hs.272897 Tubulin, alpha, brain-specific /FL=gb:BC004949.1	BC004949		NP_116093
209258_s_at	0.009689	Consensus includes gb:A1373676 /FEA=EST /DB_XREF=gi:4153542 /DB_XREF=est:qz53h11.x1 /CLONE=IMAGE:2030661 /UG=Hs.24485 chondroitin sulfate proteoglycan 6 (bamacan) /FL=gb:AF020043.1 gb:NM_005445.1 gb:AF067163.1	NM_005445		NP_005436
209259_s_at	0.02145	gb:AF020043.1 /DEF=Homo sapiens chromosome-associated polypeptide (HCAP) mRNA, complete cds. /FEA=mRNA /GEN=HCAP /PROD=chromosome associated polypeptide /DB_XREF=gi:3089367 /UG=Hs.24485 chondroitin sulfate proteoglycan 6 (bamacan) /FL=gb:AF020043.1 gb:NM_005445.1 gb:AF067163.1	AF020043		NP_005436
209269_s_at	6.68E-04	spleen tyrosine kinase	AW450910	Hs.74101	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209271_at	0.013223	gb:AB032251.1 /DEF=Homo sapiens BPTF mRNA for bromodomain PHD finger transcription factor, complete cds. /FEA=mRNA /GEN=BPTF /PROD=bromodomain PHD finger transcription factor /DB_XREF=gi:6683491 /UG=Hs.99872 fetal Alzheimer antigen /FL=gb:AB032251.1	AB032251		NP_004450
209273_s_at	1.41E-04	hypothetical protein MGC4276 similar to CG8198	BG387555	Hs.177776	NP_112202
209274_s_at	0.026891	gb:BC002675.1 /DEF=Homo sapiens, Similar to CG8198 gene product, clone MGC:4276, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG8198 gene product /DB_XREF=gi:12803678 /UG=Hs.177776 hypothetical protein MGC4276 similar to CG8198 /FL=gb:AF284752.1 gb:BC002675.1	BC002675		NP_112202
209276_s_at	3.26E-04	gb:AF162769.1 /DEF=Homo sapiens thioltransferase mRNA, complete cds. /FEA=mRNA /PROD=thioltransferase /DB_XREF=gi:5442445 /UG=Hs.28988 glutaredoxin (thioltransferase) /FL=gb:BC005304.1 gb:AF162769.1 gb:D21238.1	AF162769		NP_002055
209280_at	0.002864	gb:U58856.1 /DEF=Human chromosome 17 unknown product mRNA, complete cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:2209262 /UG=Hs.7835 endocytic receptor (macrophage mannose receptor family) /FL=gb:U58856.1 gb:AB014609.1 gb:AF134838.1 gb:NM_006039.1 gb:AF107292.1	U58856		NP_006030

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209281_s_at	5.03E-04	gb:M95541.1 /DEF=Homo sapiens adenosine triphosphatase mRNA, complete cds. /FEA=mRNA /PROD=adenosine triphosphatase /DB_XREF=gi:184269 /UG=Hs.78546 ATPase, Ca++ transporting, plasma membrane 1 /FL=gb:M95541.1 gb:NM_001682.1 gb:J04027.1	M95541		NP_001673
209284_s_at	1.26E-04	Consensus includes gb:AI922509 /FEA=EST /DB_XREF=gi:5658473 /DB_XREF=est:wh13g11.x1 /CLONE=IMAGE:2380676 /UG=Hs.23440 KIAA1105 protein /FL=gb:AF180425.2	AF180425		NP_056039
209285_s_at	0.00161	Consensus includes gb:N38985 /FEA=EST /DB_XREF=gi:1162192 /DB_XREF=est:yv21f02.s1 /CLONE=IMAGE:243387 /UG=Hs.23440 KIAA1105 protein /FL=gb:AF180425.2	AF180425		NP_056039
209286_at	0.001116	CDC42 effector protein (Rho GTPase binding) 3	AI754416	Hs.260024	
209287_s_at	0.02145	gb:AF104857.1 /DEF=Homo sapiens Cdc42 effector protein 3 mRNA, complete cds. /FEA=mRNA /PROD=Cdc42 effector protein 3 /DB_XREF=gi:4324453 /UG=Hs.260024 Cdc42 effector protein 3 /FL=gb:AF094521.1 gb:AF104857.1 gb:NM_006449.1 gb:AF164118.1 gb:AL136842.1	AF104857		NP_006440
209288_s_at	1.26E-04	gb:AL136842.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434A0530 (from clone DKFZp434A0530); complete cds. /FEA=mRNA /GEN=DKFZp434A0530 /PROD=hypothetical protein /DB_XREF=gi:6807668 /UG=Hs.260024 Cdc42 effector protein 3 /FL=gb:AF094521.1 gb:AF104857.1 gb:NM_006449.1 gb:AF164118.1 gb:AL136842.1	AL136842		NP_006440

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209293_x_at	0.00884	gb:U16153.1 /DEF=Human Id-4H protein mRNA, complete cds. /FEA=mRNA /PROD=Id-4H protein /DB_XREF=gi:625095 /UG=Hs.34853 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein /FL=gb:NM_001546.1 gb:U16153.1 gb:U28368.1	U16153		NP_001537
209295_at	5.03E-04	gb:AF016266.1 /DEF=Homo sapiens TRAIL receptor 2 mRNA, complete cds. /FEA=mRNA /PROD=TRAIL receptor 2 /DB_XREF=gi:2529562 /UG=Hs.51233 tumor necrosis factor receptor superfamily, member 10b /FL=gb:BC001281.1 gb:AF018658.1 gb:AF016849.1 gb:AF016266.1	AF016266		NP_671716
209296_at	0.00228	gb:AF136972.1 /DEF=Homo sapiens protein phosphatase 2C-like protein mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 2C-like protein /DB_XREF=gi:12239323 /UG=Hs.5687 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform /FL=gb:AF136972.1 gb:AF294792.1	AF136972		NP_808908
209300_s_at	0.005852	gb:BC002888.1 /DEF=Homo sapiens, Similar to DKFZP566B183 protein, clone MGC:10356, mRNA, complete cds. /FEA=mRNA /PROD=Similar to DKFZP566B183 protein /DB_XREF=gi:12804070 /UG=Hs.12305 DKFZP566B183 protein /FL=gb:BC002888.1	BC002888		NP_056324

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209304_x_at	3.55E-04	gb:AF087853.1 /DEF=Homo sapiens growth arrest and DNA damage inducible protein beta (GADD45B) mRNA, complete cds. /FEA=mRNA /GEN=GADD45B /PROD=growth arrest and DNA damage inducible proteinbeta /DB_XREF=gi:12061050 /UG=Hs.110571 growth arrest and DNA-damage-inducible, beta /FL=gb:AF087853.1 gb:AF078077.1	AF087853		NP_056490
209308_s_at	1.26E-04	gb:BC002461.1 /DEF=Homo sapiens, BCL2adenovirus E1B 19kD-interacting protein 2, clone MGC:1529, mRNA, complete cds. /FEA=mRNA /PROD=BCL2adenovirus E1B 19kD-interacting protein 2 /DB_XREF=gi:12803290 /UG=Hs.155596 BCL2adenovirus E1B 19kD-interacting protein 2 /FL=gb:BC002461.1	BC002461		NP_004321
209310_s_at	0.041055	gb:U25804.1 /DEF=Human Ich-2 cysteine protease mRNA, complete cds. /FEA=mRNA /PROD=Ich-2 /DB_XREF=gi:886049 /UG=Hs.74122 caspase 4, apoptosis-related cysteine protease /FL=gb:U28976.1 gb:U28977.1 gb:U28978.1 gb:NM_001225.1 gb:U25804.1 gb:U28014.1	U25804		NP_150650
209311_at	2.05E-04	gb:D87461.1 /DEF=Human mRNA for KIAA0271 gene, complete cds. /FEA=mRNA /GEN=KIAA0271 /DB_XREF=gi:1944417 /UG=Hs.75244 BCL2-like 2 /FL=gb:U59747.1 gb:D87461.1 gb:NM_004050.1	D87461		NP_004041

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209312_x_at	0.010205	gb:U65585.1 /DEF=Homo sapiens MHC class II antigen (HLA-DRB1) mRNA, HLA-DRB1*PBL allele, complete cds. /FEA=mRNA /GEN=HLA-DRB1 /PROD=MHC class II antigen /DB_XREF=gi:5478215 /UG=Hs.180255 major histocompatibility complex, class II, DR beta 1 /FL=gb:M33600.1 gb:NM_002124.1 gb:M28583.1 gb:U65585.1	U65585		NP_002115
209326_at	0.00161	gb:D84454.1 /DEF=Human mRNA for UDP-galactose translocator, complete cds. /FEA=mRNA /PROD=UDP-galactose translocator /DB_XREF=gi:1526437 /UG=Hs.21899 solute carrier family 35 (UDP-galactose transporter), member 2 /FL=gb:D84454.1	D84454		NP_005651
209333_at	6.68E-04	gb:AB018265.1 /DEF=Homo sapiens mRNA for KIAA0722 protein, complete cds. /FEA=mRNA /GEN=KIAA0722 /PROD=KIAA0722 protein /DB_XREF=gi:3882164 /UG=Hs.47061 unc-51 (C. elegans)-like kinase 1 /FL=gb:AF045458.1 gb:AB018265.1 gb:NM_003565.1	AB018265		NP_003556
209338_at	1.26E-04	gb:U03494.1 /DEF=Human transcription factor LSF mRNA, complete cds. /FEA=mRNA /PROD=transcription factor LSF /DB_XREF=gi:476098 /UG=Hs.154970 transcription factor CP2 /FL=gb:BC003634.1 gb:U03494.1	U03494		NP_005644
209339_at	3.56E-04	gb:U76248.1 /DEF=Human hSIAH2 mRNA, complete cds. /FEA=mRNA /PROD=hSIAH2 /DB_XREF=gi:2673967 /UG=Hs.20191 seven in absentia (Drosophila) homolog 2 /FL=gb:U76248.1 gb:NM_005067.1	U76248		NP_005058

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209344_at	0.041055	gb:BC002827.1 /DEF=Homo sapiens, tropomyosin 4, clone MGC:3641, mRNA, complete cds. /FEA=mRNA /PROD=tropomyosin 4 /DB_XREF=gi:12803958 /UG=Hs.250641 tropomyosin 4 /FL=gb:BC002827.1 gb:NM_003290.1	BC002827		NP_003281
209357_at	1.26E-04	gb:AF109161.1 /DEF=Homo sapiens p35srj (MRG1) mRNA, complete cds. /FEA=mRNA /GEN=MRG1 /PROD=p35srj /DB_XREF=gi:4193945 /UG=Hs.82071 Cbpp300-interacting transactivator, with GluAsp-rich carboxy-terminal domain, 2 /FL=gb:BC004377.1 gb:AF109161.1	AF109161		NP_006070
209367_at	0.00228	gb:AB002559.1 /DEF=Homo sapiens mRNA for hunc18b2, complete cds. /FEA=mRNA /PROD=hunc18b2 /DB_XREF=gi:1944129 /UG=Hs.90535 syntaxin binding protein 2 /FL=gb:BC002869.1 gb:U63533.1 gb:AB002559.1 gb:NM_006949.1	AB002559		NP_008880
209369_at	1.26E-04	gb:M63310.1 /DEF=Human 1,2-cyclic-inositol-phosphate phosphodiesterase (ANX3) mRNA, complete cds. /FEA=mRNA /GEN=ANX3 /PROD=1,2-cyclic-inositol-phosphate phosphodiesterase /DB_XREF=gi:178696 /UG=Hs.1378 annexin A3 /FL=gb:BC000871.1 gb:M63310.1 gb:M20560.1 gb:NM_005139.1	M63310		NP_005130

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209375_at	0.010205	gb:D21089.1 /DEF=Human mRNA for XP-C repair complementing protein (p125), complete cds. /FEA=mRNA /GEN=XPCC /PROD=XP-C repair complementing protein (p125) /DB_XREF=gi:475156 /UG=Hs.320 xeroderma pigmentosum, complementation group C /FL=gb:D21089.1 gb:NM_004628.1	D21089		NP_004619
209376_x_at	6.68E-04	splicing factor, arginine/serine-rich 2, interacting protein	AW084759	Hs.51957	NP_004710
209380_s_at	0.00161	gb:AF146074.1 /DEF=Homo sapiens ABC protein mRNA, complete cds. /FEA=mRNA /PROD=ABC protein /DB_XREF=gi:5006890 /UG=Hs.108660 ATP-binding cassette, sub-family C (CFTRMRP), member 5 /FL=gb:AF104942.1 gb:AB019002.1 gb:AF146074.1 gb:NM_005688.1 gb:U83661.2	AF146074		NP_005679
209382_at	0.016934	gb:U93867.1 /DEF=Human RNA polymerase III subunit (RPC62) mRNA, complete cds. /FEA=mRNA /GEN=RPC62 /PROD=RNA polymerase III subunit /DB_XREF=gi:2228747 /UG=Hs.250745 polymerase (RNA) III (DNA directed) (62kD) /FL=gb:BC002586.1 gb:NM_006468.2 gb:U93867.1	U93867		NP_006459
209384_at	5.03E-04	proline synthetase co-transcribed homolog (bacterial)	AA176833	Hs.301959	NP_009129
209388_at	0.004341	gb:BC000927.1 /DEF=Homo sapiens, Similar to poly (A) polymerase, clone MGC:5378, mRNA, complete cds. /FEA=mRNA /PROD=Similar to poly (A) polymerase /DB_XREF=gi:12654216 /UG=Hs.49007 poly(A) polymerase alpha /FL=gb:BC000927.1	BC000927		NP_116021

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209390_at	0.016934	gb:AF013168.1 /DEF=Homo sapiens hamartin (TSC1) mRNA, complete cds. /FEA=mRNA /GEN=TSC1 /PROD=hamartin /DB_XREF=gi:2331280 /UG=Hs.79393 tuberous sclerosis 1 (hamartin) /FL=gb:AF013168.1 gb:NM_000368.1	AF013168		NP_000359
209392_at	0.008237	gb:L35594.1 /DEF=Human autotaxin mRNA, complete cds. /FEA=mRNA /PROD=autotaxin /DB_XREF=gi:537905 /UG=Hs.174185 ectonucleotide pyrophosphatase phosphodiesterase 2 (autotaxin) /FL=gb:L35594.1	L35594		NP_006200
209393_s_at	1.26E-04	gb:AF047695.1 /DEF=Homo sapiens cap-binding protein 4EHP mRNA, complete cds. /FEA=mRNA /PROD=cap-binding protein 4EHP /DB_XREF=gi:3172338 /UG=Hs.19122 eukaryotic translation initiation factor 4E-like 3 /FL=gb:BC005392.1 gb:AF047695.1 gb:AF068117.1 gb:AF038957.1 gb:NM_004846.1	AF047695		NP_004837
209397_at	1.26E-04	gb:BC000147.1 /DEF=Homo sapiens, Similar to malic enzyme 2, NAD(+)-dependent, mitochondrial, clone MGC:5187, mRNA, complete cds. /FEA=mRNA /PROD=Similar to malic enzyme 2, NAD(+)-dependent, mitochondrial /DB_XREF=gi:12652790 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:BC000147.1	BC000147		NP_002387

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209403_at	0.026891	gb:AL136860.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2235 (from clone DKFZp434P2235); complete cds. /FEA=mRNA /GEN=DKFZp434P2235 /PROD=hypothetical protein /DB_XREF=gi:12053224 /UG=Hs.105891 Homo sapiens mRNA; cDNA DKFZp434P2235 (from clone DKFZp434P2235); complete cds /FL=gb:AL136860.1	AL136860		NP_115634
209404_s_at	2.05E-04	gb:AF151867.1 /DEF=Homo sapiens CGI-109 protein mRNA, complete cds. /FEA=mRNA /PROD=CGI-109 protein /DB_XREF=gi:4929686 /UG=Hs.278391 CGI-109 protein /FL=gb:AF151867.1	AF151867		
209409_at	1.26E-04	gb:D86962.1 /DEF=Human mRNA for KIAA0207 gene, complete cds. /FEA=mRNA /GEN=KIAA0207 /DB_XREF=gi:1503997 /UG=Hs.81875 growth factor receptor-bound protein 10 /FL=gb:D86962.1 gb:AF000017.1	D86962		NP_005302
209411_s_at	0.002418	golgi associated, gamma adaptin ear containing, ARF binding protein 3	AW008018	Hs.87726	NP_619525
209418_s_at	0.001116	gb:BC003615.1 /DEF=Homo sapiens, gene from NF2meningioma region of 22q12, clone MGC:1540, mRNA, complete cds. /FEA=mRNA /PROD=gene from NF2meningioma region of 22q12 /DB_XREF=gi:13177658 /UG=Hs.75361 gene from NF2meningioma region of 22q12 /FL=gb:BC003615.1 gb:NM_003678.1 gb:AB023200.1	BC003615		NP_003669

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209424_s_at	0.003817	Consensus includes gb:AI796120 /FEA=EST /DB_XREF=gi:5361583 /DB_XREF=est:wh42f03.x1 /CLONE=IMAGE:2383421 /UG=Hs.128749 alpha-methylacyl-CoA racemase /FL=gb:AF047020.1 gb:AF158378.1 gb:NM_014324.1	NM_014324		NP_055139
209426_s_at	0.01159	gb:AF047020.1 /DEF=Homo sapiens alpha-methylacyl-CoA racemase mRNA, complete cds. /FEA=mRNA /PROD=alpha-methylacyl-CoA racemase /DB_XREF=gi:4204096 /UG=Hs.128749 alpha-methylacyl-CoA racemase /FL=gb:AF047020.1 gb:AF158378.1 gb:NM_014324.1	AF047020		NP_055139
209429_x_at	0.016686	gb:AF112207.1 /DEF=Homo sapiens translation initiation factor eIF-2b delta subunit mRNA, complete cds. /FEA=mRNA /PROD=translation initiation factor eIF-2b deltasubunit /DB_XREF=gi:6563201 /UG=Hs.169474 DKFZP586J0119 protein /FL=gb:BC001870.1 gb:AF112207.1	AF112207		NP_056451
209438_at	0.004341	Consensus includes gb:AL096700 /DEF=Human DNA sequence from clone RP3-499B10 on chromosome Xp22.31-22.13. Contains the 3 end of the gene for serinethreonine protein phosphatase PPEF, the 3 end of the PHKA2 gene for liver phosphorylase kinase alpha 2, ESTs, STSs and GSSs /FEA=mRNA_2 /DB_XREF=gi:7077104 /UG=Hs.54941 phosphorylase kinase, alpha 2 (liver) /FL=gb:D38616.1 gb:NM_000292.1	AL096700		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209439_s_at	5.03E-04	gb:D38616.1 /DEF=Human mRNA for phosphorylase kinase alpha subunit, complete cds. /FEA=mRNA /PROD=phosphorylase kinase alpha subunit /DB_XREF=gi:1304117 /UG=Hs.54941 phosphorylase kinase, alpha 2 (liver) /FL=gb:D38616.1 gb:NM_000292.1	D38616		NP_000283
209447_at	0.005852	gb:AF043290.1 /DEF=Homo sapiens lymphocyte membrane associated protein (8B7) mRNA, complete cds. /FEA=mRNA /GEN=8B7 /PROD=lymphocyte membrane associated protein /DB_XREF=gi:2895592 /UG=Hs.8182 synaptic nuclei expressed gene 1b /FL=gb:AF043290.1	AF043290		NP_598411
209448_at	1.26E-04	gb:BC002439.1 /DEF=Homo sapiens, Tat-interacting protein ¹ (30kD), clone MGC:1318, mRNA, complete cds. /FEA=mRNA /PROD=Tat-interacting protein (30kD) /DB_XREF=gi:12803250 /UG=Hs.90753 Tat-interacting protein (30kD) /FL=gb:BC002439.1 gb:AF039103.1	BC002439		NP_006401
209451_at	0.013223	gb:U59863.1 /DEF=Human TRAF-interacting protein I-TRAF mRNA, complete cds. /FEA=mRNA /PROD=I-TRAF /DB_XREF=gi:1518017 /UG=Hs.146847 TRAF family member-associated NFKB activator /FL=gb:U59863.1	U59863		NP_597841

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209452_s_at	3.26E-04	gb:AF035824.1 /DEF=Homo sapiens vesicle soluble NSF attachment protein receptor (VTI1) mRNA, complete cds. /FEA=mRNA /GEN=VTI1 /PROD=vesicle soluble NSF attachment protein receptor /DB_XREF=gi:2687399 /UG=Hs.169206 vesicle-associated soluble NSF attachment protein receptor (v-SNARE; homolog of S. cerevisiae VTI1) /FL=gb:BC003142.1 gb:AF035824.1 gb:AF060902.1 gb:NM_006370.1	AF035824		NP_006361
209453_at	0.003166	gb:M81768.1 /DEF=Human NaH antiporter (APNH1) mRNA, complete cds. /FEA=mRNA /GEN=APNH /PROD=NaH antiporter /DB_XREF=gi:178752 /UG=Hs.170222 solute carrier family 9 (sodiumhydrogen exchanger), isoform 1 (antiporter, Na+H+, amiloride sensitive) /FL=gb:M81768.1 gb:AF141350.1 gb:AF141351.1 gb:AF141352.1 gb:AF141353.1 gb:AF141354.1 gb:AF141355.1 gb:AF141356.1 gb:AF141357.1 gb:AF141358.1 gb:AF141359.1 gb:AF146430.1 gb:AF146431.1 gb:AF146432.1 gb:AF146433.1 gb:AF146434.1 gb:AF146435.1 gb:AF146436.1 gb:AF146437.1 gb:AF146438.1 gb:AF146439.1	M81768		NP_003038
209455_at	1.26E-04	F-box and WD-40 domain protein 1B	BE963245	Hs.21229	NP_387449
209457_at	0.033381	gb:U16996.1 /DEF=Human protein tyrosine phosphatase mRNA, complete cds. /FEA=mRNA /PROD=protein tyrosine phosphatase /DB_XREF=gi:642012 /UG=Hs.2128 dual specificity phosphatase 5 /FL=gb:NM_004419.2 gb:U16996.1 gb:U15932.2	U16996		NP_004410

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209458_x_at	3.26E-04	gb:AF105974.1 /DEF=Homo sapiens alpha one globin (HBA1) mRNA, complete cds. /FEA=mRNA /GEN=HBA1 /PROD=alpha one globin /DB_XREF=gi:4038449 /UG=Hs.272572 hemoglobin, alpha 2 /FL=gb:AF097635.1 gb:AF105974.1 gb:NM_000517.2	AF105974		NP_000549
209459_s_at	0.041055	gb:AF237813.1 /DEF=Homo sapiens NPD009 mRNA, complete cds. /FEA=mRNA /PROD=NPD009 /DB_XREF=gi:9963907 /UG=Hs.283675 NPD009 protein /FL=gb:NM_020686.1 gb:AF237813.1	AF237813		NP_065737
209463_s_at	0.001713	gb:D50544.1 /DEF=Human TFIID subunit p22 mRNA, complete cds. /FEA=mRNA /PROD=TFIID subunit p22 /DB_XREF=gi:1345403 /UG=Hs.82037 TATA box binding protein (TBP)-associated factor, RNA polymerase II, J, 20kD /FL=gb:D50544.1 gb:NM_005644.2	D50544		NP_005635
209467_s_at	5.03E-04	gb:BC002755.1 /DEF=Homo sapiens, Similar to MAP kinase-interacting serinethreonine kinase 1, clone MGC:3690, mRNA, complete cds. /FEA=mRNA /PROD=Similar to MAP kinase-interactingserinethreonine kinase 1 /DB_XREF=gi:12803828 /UG=Hs.5591 MAP kinase-interacting serinethreonine kinase 1 /FL=gb:BC002755.1	BC002755		NP_003675
209471_s_at	0.00228	gb:L00634.1 /DEF=Human farnesyl-protein transferase alpha-subunit mRNA, complete cds. /FEA=mRNA /PROD=farnesyl-protein transferase alpha-subunit /DB_XREF=gi:292030 /UG=Hs.138381 farnesyltransferase, CAAX box, alpha /FL=gb:L00634.1 gb:L10413.1 gb:NM_002027.1	L00634		NP_002018

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209472_at	0.007779	gb:BC000819.1 /DEF=Homo sapiens, Similar to CG6950 gene product, clone MGC:5114, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG6950 gene product /DB_XREF=gi:12654030 /UG=Hs.180378 hypothetical protein 669 /FL=gb:BC000819.1 gb:NM_019610.1	BC000819		NP_062556
209473_at	1.26E-04	ectonucleoside triphosphate diphosphohydrolase 1	AV717590	Hs.205353	NP_001767
209475_at	0.004341	gb:AF106069.1 /DEF=Homo sapiens deubiquitinating enzyme (UNPH4) mRNA, complete cds. /FEA=mRNA /GEN=UNPH4 /PROD=deubiquitinating enzyme /DB_XREF=gi:5814096 /UG=Hs.23168 ubiquitin specific protease 15 /FL=gb:AF013990.1 gb:AF106069.1	AF106069		NP_006304
209476_at	5.03E-04	Consensus includes gb:AL080080.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E1962 (from clone DKFZp564E1962); partial cds. /FEA=mRNA /GEN=DKFZp564E1962 /PROD=hypothetical protein /DB_XREF=gi:5262491 /UG=Hs.24766 thioredoxin-related transmembrane protein /FL=gb:AB048246.1	AL080080		NP_110382
209479_at	1.26E-04	gb:BC000758.1 /DEF=Homo sapiens, clone MGC:2698, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:2698) /DB_XREF=gi:12653928 /UG=Hs.44468 DKFZP586D0623 protein /FL=gb:BC000758.1	BC000758		NP_056254

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209480_at	0.022752	gb:M16276.1 /DEF=Human MHC class II HLA-DR2-Dw12 mRNA DQw1-beta, complete cds. /FEA=mRNA /GEN=HLA-DRB2 /DB_XREF=gi:188397 /UG=Hs.73931 major histocompatibility complex, class II, DQ beta 1 /FL=gb:M60028.1 gb:M17564.1 gb:M81140.1 gb:M81141.1 gb:M16276.1 gb:NM_002123.1	M16276		NP_002114
209481_at	0.005852	gb:AF226044.1 /DEF=Homo sapiens HSNFRK (HSNFRK) mRNA, complete cds. /FEA=mRNA /GEN=HSNFRK /PROD=HSNFRK /DB_XREF=gi:9295326 /UG=Hs.79025 KIAA0096 protein /FL=gb:AF226044.1	AF226044		NP_060189
209482_at	0.005339	gb:BC001430.1 /DEF=Homo sapiens, POP7 (processing of precursor, S. cerevisiae) homolog, clone MGC:1986, mRNA, complete cds. /FEA=mRNA /PROD=POP7 (processing of precursor, S. cerevisiae)homolog /DB_XREF=gi:12655150 /UG=Hs.18747 POP7 (processing of precursor, S. cerevisiae) homolog /FL=gb:BC001430.1 gb:U94316.1 gb:NM_005837.1	BC001430		NP_005828
209497_s_at	0.02145	gb:BC003503.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 4921506I22 gene, clone MGC:10380, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 4921506I22 gene /DB_XREF=gi:13097557 /UG=Hs.49994 Homo sapiens, clone MGC:10871, mRNA, complete cds /FL=gb:BC003503.1 gb:BC004951.1	BC003503		NP_113680

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209498_at	3.56E-04	Consensus includes gb:X16354.1 /DEF=Human mRNA for transmembrane carcinoembryonic antigen BGPα (formerly TM1-CEA). /FEA=mRNA /PROD=TM1-CEA preprotein /DB_XREF=gi:37197 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:J03858.1	X16354		NP_001703
209500_x_at	0.00228	gb:AF114012.1 /DEF=Homo sapiens tumor necrosis factor-related death ligand-1β mRNA, complete cds. /FEA=mRNA /PROD=tumor necrosis factor-related deathligand-1β /DB_XREF=gi:7328555 /UG=Hs.54673 tumor necrosis factor (ligand) superfamily, member 13 /FL=gb:AF114012.1	AF114012		NP_742086
209510_at	0.007779	gb:AF064801.1 /DEF=Homo sapiens multiple membrane spanning receptor TRC8 (TRC8) mRNA, complete cds. /FEA=mRNA /GEN=TRC8 /PROD=multiple membrane spanning receptor TRC8 /DB_XREF=gi:3395786 /UG=Hs.28285 patched related protein translocated in renal cancer /FL=gb:AF064801.1 gb:NM_007218.1	AF064801		NP_009149
209512_at	3.26E-04	gb:BC004331.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 2610207I16 gene /DB_XREF=gi:13279253 /UG=Hs.47986 Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds /FL=gb:BC004331.1	BC004331		NP_115679

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209513_s_at	3.26E-04	gb:BC004331.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 2610207I16 gene /DB_XREF=gi:13279253 /UG=Hs.47986 Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds /FL=gb:BC004331.1	BC004331		NP_115679
209514_s_at	0.00161	RAB27A, member RAS oncogene family	BE502030	Hs.50477	NP_004571
209515_s_at	2.05E-04	gb:U38654.3 /DEF=Homo sapiens Rab27a mRNA, complete cds. /FEA=mRNA /PROD=Rab27a /DB_XREF=gi:4887230 /UG=Hs.50477 RAB27A, member RAS oncogene family /FL=gb:U38654.3	U38654		NP_004571
209517_s_at	3.26E-04	gb:AB020982.1 /DEF=Homo sapiens ASH2L mRNA, complete cds, similar to Drosophila ash2 sequence. /FEA=mRNA /GEN=ASH2L /DB_XREF=gi:4417209 /UG=Hs.6856 ash2 (absent, small, or homeotic, Drosophila, homolog)-like /FL=gb:AF056718.1 gb:AB020982.1 gb:NM_004674.1	AB020982		NP_004665
209520_s_at	0.007779	gb:BC001450.1 /DEF=Homo sapiens, nuclear cap binding protein subunit 1, 80kD, clone MGC:2087, mRNA, complete cds. /FEA=mRNA /PROD=nuclear cap binding protein subunit 1, 80kD /DB_XREF=gi:12655186 /UG=Hs.89563 nuclear cap binding protein subunit 1, 80kD /FL=gb:BC001450.1 gb:D32002.1 gb:NM_002486.1	BC001450		NP_002477

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209525_at	0.012818	Consensus includes gb:BG285017 /FEA=EST /DB_XREF=gi:13036553 /DB_XREF=est:602409285F1 /CLONE=IMAGE:4538229 /UG=Hs.127842 CGI-142 /FL=gb:AF151900.1 gb:AB029156.1 gb:NM_016073.1	NM_016073		NP_057157
209539_at	0.013223	Consensus includes gb:D25304.1 /DEF=Human mRNA for KIAA0006 gene, partial cds. /FEA=mRNA /GEN=KIAA0006 /DB_XREF=gi:435445 /UG=Hs.79307 RacCdc42 guanine exchange factor (GEF) 6 /FL=gb:D13631.1	D25304		NP_004831
209549_s_at	7.58E-04	gb:BC001121.1 /DEF=Homo sapiens, Similar to deoxyguanosine kinase, clone MGC:2111, mRNA, complete cds. /FEA=mRNA /PROD=Similar to deoxyguanosine kinase /DB_XREF=gi:12654572 /UG=Hs.326494 Homo sapiens, Similar to deoxyguanosine kinase, clone MGC:2111, mRNA, complete cds /FL=gb:BC001121.1	BC001121		NP_550440
209551_at	3.26E-04	gb:BC004875.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 2310034L04 gene, clone MGC:11061, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 2310034L04 gene /DB_XREF=gi:13436109 /UG=Hs.66309 Homo sapiens, Similar to RIKEN cDNA 2310034L04 gene, clone MGC:11061, mRNA, complete cds /FL=gb:BC004875.1	BC004875		NP_115688

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209553_at	1.26E-04	gb:BC001001.2 /DEF=Homo sapiens, Similar to hBKLF for basic kruppel like factor, clone MGC:5362, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hBKLF for basic kruppel like factor /DB_XREF=gi:12803042 /UG=Hs.7316 KIAA0804 protein /FL=gb:BC001001.2	BC001001		
209555_s_at	0.00161	gb:M98399.1 /DEF=Human antigen CD36 (clone 21) mRNA, complete cds. /FEA=mRNA /GEN=CD36 /PROD=antigen CD36 /DB_XREF=gi:180112 /UG=Hs.75613 CD36 antigen (collagen type I receptor, thrombospondin receptor) /FL=gb:M98399.1	M98399		NP_000063
209561_at	0.020456	gb:L38969.1 /DEF=Homo sapiens thrombospondin 3 (THBS3) mRNA, complete cds. /FEA=mRNA /GEN=THBS3 /PROD=thrombospondin 3 /DB_XREF=gi:886298 /UG=Hs.169875 thrombospondin 3 /FL=gb:NM_007112.1 gb:L38969.1	L38969		NP_009043
209566_at	0.001116	Consensus includes gb:AL080184.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O071 (from clone DKFZp434O071). /FEA=mRNA /DB_XREF=gi:5262661 /UG=Hs.7089 insulin induced protein 2 /FL=gb:AF125392.1	AL080184		NP_057217
209574_s_at	0.00161	chromosome 18 open reading frame 1	AI349506	Hs.153498	NP_004329

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209575_at	0.010205	gb:BC001903.1 /DEF=Homo sapiens, Similar to interleukin 10 receptor, beta, clone MGC:2210, mRNA, complete cds. /FEA=mRNA /PROD=Similar to interleukin 10 receptor, beta /DB_XREF=gi:12804902 /UG=Hs.173936 interleukin 10 receptor, beta /FL=gb:BC001903.1 gb:NM_000628.1	BC001903		NP_000619
209579_s_at	0.00161	methyl-CpG binding domain protein 4	AL556619	Hs.35947	NP_003916
209580_s_at	0.004341	gb:AF114784.1 /DEF=Homo sapiens methyl-CpG binding endonuclease (MED1) mRNA, complete cds. /FEA=mRNA /GEN=MED1 /PROD=methyl-CpG binding endonuclease /DB_XREF=gi:4539758 /UG=Hs.35947 methyl-CpG binding domain protein 4 /FL=gb:AF072250.1 gb:NM_003925.1 gb:AF114784.1	AF114784		NP_003916
209593_s_at	0.00228	gb:AF317129.1 /DEF=Homo sapiens FKSG18 (FKSG18) mRNA, complete cds. /FEA=mRNA /GEN=FKSG18 /PROD=FKSG18 /DB_XREF=gi:12276123 /UG=Hs.252682 torsin family 1, member B (torsin B) /FL=gb:AF317129.1	AF317129		NP_055321
209595_at	0.002313	gb:BC001771.1 /DEF=Homo sapiens, general transcription factor IIF, polypeptide 2 (30kD subunit), clone MGC:1502, mRNA, complete cds. /FEA=mRNA /PROD=general transcription factor IIF, polypeptide 2(30kD subunit) /DB_XREF=gi:12804688 /UG=Hs.58593 general transcription factor IIF, polypeptide 2 (30kD subunit) /FL=gb:BC001771.1 gb:NM_004128.1	BC001771		NP_004119
209599_s_at	0.045257	TcD37 homolog	AI762105	Hs.332683	NP_067045

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209604_s_at	0.00228	gb:BC003070.1 /DEF=Homo sapiens, GATA-binding protein 3, clone MGC:2346, mRNA, complete cds. /FEA=mRNA /PROD=GATA-binding protein 3 /DB_XREF=gi:13111765 /UG=Hs.169946 GATA-binding protein 3 /FL=gb:BC003070.1 gb:M69106.1 gb:NM_002051.1	BC003070		NP_002042
209605_at	0.007943	gb:D87292.1 /DEF=Homo sapiens mRNA for rhodanese, complete cds. /FEA=mRNA /PROD=rhodanese /DB_XREF=gi:1877030 /UG=Hs.248267 thiosulfate sulfurtransferase (rhodanese) /FL=gb:D87292.1	D87292		NP_003303
209610_s_at	5.64E-04	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	BF340083	Hs.323878	NP_003029
209615_s_at	0.004341	gb:U51120.1 /DEF=Human protein kinase PAK1 mRNA, complete cds. /FEA=mRNA /PROD=PAK1 /DB_XREF=gi:1256421 /UG=Hs.62402 p21Cdc42Rac1-activated kinase 1 (yeast Ste20-related) /FL=gb:U51120.1 gb:NM_002576.2 gb:U24152.1	U51120		NP_002567
209619_at	3.26E-04	gb:K01144.1 /DEF=Human major histocompatibility class II antigen gamma chain mRNA, complete cds. /FEA=mRNA /PROD=class II antigen gamma chain /DB_XREF=gi:188469 /UG=Hs.84298 CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) /FL=gb:NM_004355.1 gb:K01144.1	K01144		NP_004346

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209620_s_at	0.022752	gb:AB005289.1 /DEF=Homo sapiens mRNA for ABC transporter 7 protein, complete cds. /FEA=mRNA /GEN=hABC7 /PROD=ABC transporter 7 protein /DB_XREF=gi:3228278 /UG=Hs.125856 ATP-binding cassette, sub-family B (MDRTAP), member 7 /FL=gb:AB005289.1 gb:AF038950.1 gb:AF133659.1 gb:NM_004299.2	AB005289		NP_004290
209628_at	0.007779	Consensus includes gb:AK023289.1 /DEF=Homo sapiens cDNA FLJ13227 fis, clone OVARC1000071, weakly similar to Homo sapiens NTF2-related export protein NXT1 (NXT1) mRNA. /FEA=mRNA /DB_XREF=gi:10435160 /UG=Hs.25010 hypothetical protein P15-2 /FL=gb:AF246127.1 gb:NM_018698.1 gb:AF201942.1	AK023289		NP_061168
209630_s_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AL043967	Hs.350947	
209640_at	3.26E-04	gb:M79462.1 /DEF=Human PML-1 mRNA, complete CDS. /FEA=mRNA /GEN=PML /DB_XREF=gi:190116 /UG=Hs.89633 promyelocytic leukemia /FL=gb:M79462.1	M79462		NP_150253
209643_s_at	0.002537	gb:AF033850.1 /DEF=Homo sapiens phospholipase D2 (PLD2) mRNA, complete cds. /FEA=mRNA /GEN=PLD2 /PROD=phospholipase D2 /DB_XREF=gi:2645857 /UG=Hs.104519 phospholipase D2 /FL=gb:AF033850.1 gb:AF038440.1 gb:AF035483.1 gb:NM_002663.1	AF033850		NP_002654

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209649_at	0.005339	gb:AL133600.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564C047 (from clone DKFZp564C047); complete cds. /FEA=mRNA /GEN=DKFZp564C047 /PROD=hypothetical protein /DB_XREF=gi:6599205 /UG=Hs.17200 STAM-like protein containing SH3 and ITAM domains 2 /FL=gb:AF042273.1 gb:AL133600.1	AL133600		NP_005834
209653_at	6.93E-04	gb:U93240.1 /DEF=Human importin alpha 3 mRNA, complete cds. /FEA=mRNA /PROD=importin alpha 3 /DB_XREF=gi:1928974 /UG=Hs.302499 karyopherin alpha 4 (importin alpha 3) /FL=gb:AB002533.2 gb:NM_002268.2 gb:U93240.1	U93240		NP_002259
209665_at	0.041055	gb:AF040704.1 /DEF=Homo sapiens putative tumor suppressor protein (101F6) mRNA, complete cds. /FEA=mRNA /GEN=101F6 /PROD=putative tumor suppressor protein /DB_XREF=gi:2935319 /UG=Hs.149443 putative tumor suppressor /FL=gb:AF040704.1 gb:NM_007022.1	AF040704		NP_008953
209666_s_at	2.05E-04	gb:AF080157.1 /DEF=Homo sapiens Ikb kinase-a (IKK-alpha) mRNA, complete cds. /FEA=mRNA /GEN=IKK-alpha /PROD=Ikb kinase-a /DB_XREF=gi:4185272 /UG=Hs.198998 conserved helix-loop-helix ubiquitous kinase /FL=gb:U22512.1 gb:AF012890.1 gb:AF009225.1 gb:AF080157.1 gb:NM_001278.1	AF080157		NP_001269
209667_at	1.32E-04	carboxylesterase 2 (intestine, liver)	BF033242	Hs.282975	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209670_at	0.005852	gb:M12959.1 /DEF=Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds. /FEA=mRNA /GEN=TCRA /DB_XREF=gi:338734 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:M12959.1 gb:M12423.1	M12959		
209671_x_at	0.001618	gb:M12423.1 /DEF=Human T-cell receptor active alpha-chain mRNA, complete cds, from Jurkat T-cell line. /FEA=mRNA /GEN=TCRA /DB_XREF=gi:338738 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:M12959.1 gb:M12423.1	M12423		
209678_s_at	2.05E-04	gb:L18964.1 /DEF=Human protein kinase C iota isoform (PRKCI) mRNA, complete cds. /FEA=mRNA /GEN=PRKCI /PROD=protein kinase C iota /DB_XREF=gi:432273 /UG=Hs.1904 protein kinase C, iota /FL=gb:L18964.1 gb:NM_002740.1 gb:L33881.1	L18964		NP_002731
209681_at	0.004733	gb:AF153330.1 /DEF=Homo sapiens thiamine carrier 1 (TC1) mRNA, complete cds. /FEA=mRNA /GEN=TC1 /PROD=thiamine carrier 1 /DB_XREF=gi:5453325 /UG=Hs.30246 solute carrier family 19 (thiamine transporter), member 2 /FL=gb:AF153330.1 gb:AF135488.1 gb:AF160812.1	AF153330		NP_008927
209682_at	5.03E-04	gb:U26710.1 /DEF=Human cbl-b mRNA, complete cds. /FEA=mRNA /PROD=cbl-b /DB_XREF=gi:862406 /UG=Hs.3144 Cas-Br-M (murine) ectropic retroviral transforming sequence b /FL=gb:U26710.1	U26710		NP_733762

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209685_s_at	0.00228	gb:M13975.1 /DEF=Homo sapiens protein kinase C beta-II type (PRKCB1) mRNA, complete cds. /FEA=mRNA /GEN=PRKCB1 /PROD=protein kinase C beta-II type /DB_XREF=gi:189968 /UG=Hs.77202 protein kinase C, beta 1 /FL=gb:M13975.1	M13975		NP_002729
209688_s_at	0.00228	gb:BC005078.1 /DEF=Homo sapiens, clone MGC:13033, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13033) /DB_XREF=gi:13477224 /UG=Hs.26118 Homo sapiens clone 24766 mRNA sequence /FL=gb:BC005078.1	BC005078		NP_113635
209689_at	0.011788	gb:BC005078.1 /DEF=Homo sapiens, clone MGC:13033, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13033) /DB_XREF=gi:13477224 /UG=Hs.26118 Homo sapiens clone 24766 mRNA sequence /FL=gb:BC005078.1	BC005078		NP_113635
209694_at	0.045608	gb:M97655.1 /DEF=Human 6-pyruvoyltetrahydropterin synthase (PTS) mRNA, complete cds. /FEA=mRNA /GEN=PTS /PROD=6-pyruvoyltetrahydropterin synthase /DB_XREF=gi:306438 /UG=Hs.366 6-pyruvoyltetrahydropterin synthase /FL=gb:M97655.1 gb:NM_000317.1 gb:D17400.1	M97655		NP_000308
209707_at	1.26E-04	gb:AF022913.1 /DEF=Homo sapiens GPI transamidase mRNA, complete cds. /FEA=mRNA /PROD=GPI transamidase /DB_XREF=gi:2558890 /UG=Hs.62187 phosphatidylinositol glycan, class K /FL=gb:AF022913.1	AF022913		NP_005473
209711_at	0.003171	UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter	N80922	Hs.82635	NP_055954

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209732_at	0.02145	gb:BC005254.1 /DEF=Homo sapiens, Similar to C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced), clone MGC:12289, mRNA, complete cds. /FEA=mRNA /PROD=Similar to C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced) /DB_XREF=gi:13528920 /UG=Hs.85201 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced) /FL=gb:BC005254.1 gb:AB015628.1 gb:NM_005127.1	BC005254		NP_005118
209734_at	2.05E-04	gb:BC001604.1 /DEF=Homo sapiens, hematopoietic protein 1, clone MGC:2204, mRNA, complete cds. /FEA=mRNA /PROD=hematopoietic protein 1 /DB_XREF=gi:12804404 /UG=Hs.132834 hematopoietic protein 1 /FL=gb:BC001604.1 gb:M58285.1 gb:NM_005337.1	BC001604		NP_005328
209738_x_at	0.022758	gb:M31125.1 /DEF=Human pregnancy-specific-beta 1-glycoprotein (PS1) mRNA, complete cds, clone hPS12. /FEA=mRNA /DB_XREF=gi:190619 /UG=Hs.252097 pregnancy specific beta-1-glycoprotein 6 /FL=gb:M31125.1	M31125		NP_002773
209748_at	0.026891	gb:AB029006.1 /DEF=Homo sapiens mRNA for KIAA1083 protein, complete cds. /FEA=mRNA /GEN=KIAA1083 /PROD=KIAA1083 protein /DB_XREF=gi:5689502 /UG=Hs.26334 spastic paraplegia 4 (autosomal dominant; spastin) /FL=gb:AB029006.1	AB029006		NP_055761
209753_s_at	0.001116	thymopoietin	BG391171	Hs.11355	NP_003267

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209754_s_at	0.011626	gb:AF113682.1 /DEF=Homo sapiens clone FLB3436 PRO0868 mRNA, complete cds. /FEA=mRNA /PROD=PRO0868 /DB_XREF=gi:6855608 /UG=Hs.11355 thymopoietin /FL=gb:U09087.1 gb:AF113682.1	AF113682		NP_003267
209760_at	1.26E-04	gb:AL136932.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H1322 (from clone DKFZp586H1322); complete cds. /FEA=mRNA /GEN=DKFZp586H1322 /PROD=hypothetical protein /DB_XREF=gi:12053358 /UG=Hs.37892 KIAA0922 protein /FL=gb:AL136932.1	AL136932		NP_056011
209761_s_at	0.004341	Consensus includes gb:AA969194 /FEA=EST /DB_XREF=gi:3144374 /DB_XREF=est:op51c02.s1 /CLONE=IMAGE:1580354 /UG=Hs.38125 interferon-induced protein 75, 52kD /FL=gb:AF280094.1	AF280094		NP_536349
209771_x_at	1.26E-04	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	AA761181	Hs.381003	NP_037362
209772_s_at	0.004659	gb:X69397.1 /DEF=H.sapiens CD24 gene, complete CDS. /FEA=mRNA /GEN=CD24 /PROD=cell surface antigen /DB_XREF=gi:396167 /UG=Hs.286124 CD24 antigen (small cell lung carcinoma cluster 4 antigen) /FL=gb:X69397.1	X69397		NP_037362
209773_s_at	1.58E-04	gb:BC001886.1 /DEF=Homo sapiens, ribonucleotide reductase M2 polypeptide, clone MGC:1551, mRNA, complete cds. /FEA=mRNA /PROD=ribonucleotide reductase M2 polypeptide /DB_XREF=gi:12804874 /UG=Hs.75319 ribonucleotide reductase M2 polypeptide /FL=gb:BC001886.1	BC001886		NP_001025

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209788_s_at	0.001618	gb:AF183569.1 /DEF=Homo sapiens aminopeptidase PILS (APPILS) mRNA, complete cds. /FEA=mRNA /GEN=APPILS /PROD=aminopeptidase PILS /DB_XREF=gi:6642986 /UG=Hs.247043 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator /FL=gb:AF183569.1 gb:AF222340.1 gb:NM_016442.1	AF183569		NP_057526
209799_at	0.001109	gb:AF100763.1 /DEF=Homo sapiens AMP-activated kinase alpha 1 subunit mRNA, complete cds. /FEA=mRNA /PROD=AMP-activated kinase alpha 1 subunit /DB_XREF=gi:5410311 /UG=Hs.9247 protein kinase, AMP-activated, alpha 1 catalytic subunit /FL=gb:AB022017.1 gb:AF100763.1 gb:NM_006251.1	AF100763		NP_006242
209806_at	1.26E-04	gb:BC000893.1 /DEF=Homo sapiens, H2B histone family, member A, clone MGC:5132, mRNA, complete cds. /FEA=mRNA /PROD=H2B histone family, member A /DB_XREF=gi:12654150 /UG=Hs.247817 H2B histone family, member A /FL=gb:BC000893.1	BC000893		NP_542160
209819_at	0.041795	gb:AF241831.1 /DEF=Homo sapiens intracellular hyaluronan-binding protein mRNA, complete cds. /FEA=mRNA /PROD=intracellular hyaluronan-binding protein /DB_XREF=gi:7417371 /UG=Hs.301839 intracellular antigen detected by monoclonal antibody Ki-1; intracellular hyaluronan-binding protein /FL=gb:AF241831.1	AF241831		NP_055097

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209829_at	0.001116	gb:AB002384.1 /DEF=Human mRNA for KIAA0386 gene, complete cds. /FEA=mRNA /GEN=KIAA0386 /DB_XREF=gi:2224712 /UG=Hs.101359 chromosome 6 open reading frame 32 /FL=gb:AB002384.1	AB002384		NP_056948
209831_x_at	0.001198	gb:AB004574.1 /DEF=Homo sapiens mRNA for deoxyribonuclease II, complete cds. /FEA=mRNA /GEN=DNASE2 /PROD=deoxyribonuclease II /DB_XREF=gi:3184394 /UG=Hs.118243 deoxyribonuclease II, lysosomal /FL=gb:AF045937.1 gb:AF047016.1 gb:AB004574.1 gb:AF060222.1 gb:NM_001375.1	AB004574		NP_001366
209833_at	1.26E-04	gb:U79115.1 /DEF=Human death adaptor molecule RAIDD (RAIDD) mRNA, complete cds. /FEA=mRNA /GEN=RAIDD /PROD=death adaptor molecule RAIDD /DB_XREF=gi:1785556 /UG=Hs.155566 CASP2 and RIPK1 domain containing adaptor with death domain /FL=gb:U79115.1 gb:U84388.1 gb:NM_003805.1	U79115		NP_003796
209845_at	0.00161	gb:AF117233.1 /DEF=Homo sapiens znf-xp protein mRNA, complete cds. /FEA=mRNA /PROD=znf-xp protein /DB_XREF=gi:6563239 /UG=Hs.7838 makorin, ring finger protein, 1 /FL=gb:AF117233.1	AF117233		NP_038474

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209846_s_at	2.05E-04	gb:BC002832.1 /DEF=Homo sapiens, Similar to butyrophilin, subfamily 3, member A2, clone MGC:3790, mRNA, complete cds. /FEA=mRNA /PROD=Similar to butyrophilin, subfamily 3, member A2 /DB_XREF=gi:12803966 /UG=Hs.87497 butyrophilin, subfamily 3, member A2 /FL=gb:BC002832.1 gb:U90144.1 gb:U90546.1 gb:NM_007047.1	BC002832		NP_008978
209852_x_at	1.26E-04	gb:BC001423.1 /DEF=Homo sapiens, Similar to proteasome (prosome, macropain) 28 subunit, 3, clone MGC:1394, mRNA, complete cds. /FEA=mRNA /PROD=Similar to proteasome (prosome, macropain) 28subunit, 3 /DB_XREF=gi:12655138 /UG=Hs.152978 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) /FL=gb:BC001423.1 gb:BC002684.1	BC001423		NP_789839
209853_s_at	0.003741	gb:BC002684.1 /DEF=Homo sapiens, proteasome (prosome, macropain) 28 subunit, 3, clone MGC:3572, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) 28 subunit, 3 /DB_XREF=gi:12803696 /UG=Hs.152978 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) /FL=gb:BC001423.1 gb:BC002684.1	BC002684		NP_789839

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209858_x_at	0.003171	gb:BC002877.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ11585 /DB_XREF=gi:12804048 /UG=Hs.315367 Homo sapiens, Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA, complete cds /FL=gb:BC002877.1	BC002877		NP_075563
209860_s_at	0.001116	gb:J04543.1 /DEF=Human synexin mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:338243 /UG=Hs.78637 annexin A7 /FL=gb:BC002632.1 gb:J04543.1 gb:NM_001156.2	J04543		NP_004025
209864_at	0.004341	gb:AB045118.1 /DEF=Homo sapiens FRAT2 mRNA, complete cds. /FEA=mRNA /GEN=FRAT2 /PROD=FRAT2 /DB_XREF=gi:13365650 /UG=Hs.140720 GSK-3 binding protein FRAT2 /FL=gb:AB045118.1	AB045118		NP_036215
209868_s_at	1.26E-04	gb:D28482.1 /DEF=Human scr2 mRNA for RNA binding protein SCR2, complete cds. /FEA=mRNA /GEN=scr2 /PROD=SCR2 /DB_XREF=gi:520588 /UG=Hs.241567 RNA binding motif, single stranded interacting protein 1 /FL=gb:D28482.1 gb:NM_016836.1 gb:NM_016838.1 gb:NM_002897.2	D28482		NP_058523

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209882_at	3.26E-04	gb:AF084462.1 /DEF=Homo sapiens GTP-binding protein ROC1 (ROC1) mRNA, complete cds. /FEA=mRNA /GEN=ROC1 /PROD=GTP-binding protein ROC1 /DB_XREF=gi:4234917 /UG=Hs.96038 Ric (Drosophila)-like, expressed in many tissues /FL=gb:U71203.1 gb:U78165.1 gb:AF084462.1 gb:NM_006912.1	AF084462		NP_008843
209892_at	1.72E-04	Consensus includes gb:AF305083.1 /DEF=Homo sapiens alpha(1,3)-fucosyltransferase IV (FUTIV) gene, 3 UTR. /FEA=mRNA /DB_XREF=gi:11096240 /UG=Hs.2173 fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific) /FL=gb:M58596.1 gb:M58597.1 gb:NM_002033.1	AF305083		NP_002024
209894_at	0.006698	gb:U50748.1 /DEF=Homo sapiens leptin receptor short form (db) mRNA, complete cds. /FEA=mRNA /GEN=db /PROD=leptin receptor /DB_XREF=gi:3236285 /UG=Hs.226627 leptin receptor /FL=gb:U50748.1	U50748		NP_002294
209896_s_at	0.02145	gb:AF119855.1 /DEF=Homo sapiens PRO1847 mRNA, complete cds. /FEA=mRNA /PROD=PRO1847 /DB_XREF=gi:7770146 /UG=Hs.285196 hypothetical protein PRO1847 /FL=gb:AF119855.1	AF119855		
209901_x_at	7.58E-04	gb:U19713.1 /DEF=Human allograft-inflammatory factor-1 mRNA, complete cds. /FEA=mRNA /PROD=allograft-inflammatory factor-1 /DB_XREF=gi:1122908 /UG=Hs.76364 allograft inflammatory factor 1 /FL=gb:U19713.1 gb:U49392.1 gb:D86438.1 gb:NM_001623.2	U19713		NP_116573

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209906_at	2.05E-04	gb:U62027.1 /DEF=Human anaphylatoxin C3a receptor (HNFAG09) mRNA, complete cds. /FEA=mRNA /GEN=HNFAG09 /PROD=anaphylatoxin C3a receptor /DB_XREF=gi:1511643 /UG=Hs.155935 complement component 3a receptor 1 /FL=gb:U28488.1 gb:U62027.1 gb:NM_004054.1	U62027		NP_004045
209907_s_at	0.00228	gb:AF182198.1 /DEF=Homo sapiens intersectin 2 long isoform (ITSN2) mRNA, complete cds. /FEA=mRNA /GEN=ITSN2 /PROD=intersectin 2 long isoform /DB_XREF=gi:7329075 /UG=Hs.166184 intersectin 2 /FL=gb:AF182198.1	AF182198		NP_671494
209911_x_at	0.001116	gb:BC002842.1 /DEF=Homo sapiens, H2B histone family, member B, clone MGC:3802, mRNA, complete cds. /FEA=mRNA /PROD=H2B histone family, member B /DB_XREF=gi:12803984 /UG=Hs.180779 H2B histone family, member B /FL=gb:NM_021063.1 gb:BC002842.1	BC002842		NP_619790
209916_at	3.56E-04	gb:BC002477.1 /DEF=Homo sapiens, clone MGC:3090, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3090) /DB_XREF=gi:12803318 /UG=Hs.271586 hypothetical protein DKFZp762M115 /FL=gb:BC002477.1	BC002477		NP_061176
209919_x_at	2.05E-04	gb:L20490.1 /DEF=Human gamma-glutamyl transpeptidase mRNA, complete cds. /FEA=mRNA /PROD=gamma-glutamyl transpeptidase /DB_XREF=gi:306748 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:J05235.1 gb:L20490.1	L20490		NP_543029

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209930_s_at	0.026891	gb:L13974.1 /DEF=Human leucine zipper mRNA, complete cds. /FEA=mRNA /PROD=leucine zipper protein /DB_XREF=gi:506817 /UG=Hs.75643 nuclear factor (erythroid-derived 2), 45kD /FL=gb:BC005044.1 gb:L24122.1 gb:L13974.1	L13974		NP_006154
209943_at	0.00161	gb:AF176699.1 /DEF=Homo sapiens F-box protein FBL4 mRNA, complete cds. /FEA=mRNA /PROD=F-box protein FBL4 /DB_XREF=gi:6103636 /UG=Hs.49526 f-box and leucine-rich repeat protein 4 /FL=gb:AF176699.1 gb:AF199355.1 gb:NM_012160.1	AF176699		NP_036292
209944_at	3.26E-04	gb:BC000330.1 /DEF=Homo sapiens, Similar to clones 23667 and 23775 zinc finger protein, clone MGC:8520, mRNA, complete cds. /FEA=mRNA /PROD=Similar to clones 23667 and 23775 zinc fingerprotein /DB_XREF=gi:12653126 /UG=Hs.7137 clones 23667 and 23775 zinc finger protein /FL=gb:BC000330.1 gb:BC004357.1	BC000330		NP_067011
209960_at	0.001053	Consensus includes gb:X16323.1 /DEF=Human mRNA for hepatocyte growth factor (HGF). /FEA=mRNA /DB_XREF=gi:32081 /UG=Hs.809 hepatocyte growth factor (hepapoietin A; scatter factor) /FL=gb:M60718.1 gb:M29145.1 gb:M73239.1 gb:M73240.1	X16323		NP_000592

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209964_s_at	0.007779	gb:AF032105.1 /DEF=Homo sapiens ataxin-7 (SCA7) mRNA, complete cds. /FEA=mRNA /GEN=SCA7 /PROD=ataxin-7 /DB_XREF=gi:3192953 /UG=Hs.108447 spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration) /FL=gb:AF032105.1	AF032105		NP_000324
209984_at	0.003171	gb:AB037901.1 /DEF=Homo sapiens GASC-1 mRNA, complete cds. /FEA=mRNA /GEN=GASC-1 /DB_XREF=gi:10567163 /UG=Hs.149918 gene amplified in squamous cell carcinoma 1; KIAA0780 protein /FL=gb:AB037901.1	AB037901		NP_055876
209992_at	7.22E-04	gb:AB044805.1 /DEF=Homo sapiens mRNA for 6-phosphofructo-2-kinase heart isoform, complete cds. /FEA=mRNA /PROD=6-phosphofructo-2-kinase heart isoform /DB_XREF=gi:11933148 /UG=Hs.211585 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 2 /FL=gb:AB044805.1	AB044805		NP_006203
209993_at	0.023827	gb:AF016535.1 /DEF=Homo sapiens P-glycoprotein (mdr1) mRNA, complete cds. /FEA=mRNA /GEN=mdr1 /PROD=P-glycoprotein /DB_XREF=gi:2353263 /UG=Hs.21330 ATP-binding cassette, sub-family B (MDRTAP), member 1 /FL=gb:M14758.1 gb:AF016535.1 gb:NM_000927.2	AF016535		NP_000918

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209997_x_at	0.009652	gb:BC000453.1 /DEF=Homo sapiens, Similar to pericentriolar material 1, clone MGC:8458, mRNA, complete cds. /FEA=mRNA /PROD=Similar to pericentriolar material 1 /DB_XREF=gi:12653366 /UG=Hs.315478 Homo sapiens, Similar to pericentriolar material 1, clone MGC:8458, mRNA, complete cds /FL=gb:BC000453.1	BC000453		NP_006188
210004_at	1.28E-04	gb:AF035776.1 /DEF=Homo sapiens oxidized low-density lipoprotein receptor mRNA, complete cds. /FEA=mRNA /PROD=oxidized low-density lipoprotein receptor /DB_XREF=gi:3941299 /UG=Hs.77729 oxidised low density lipoprotein (lectin-like) receptor 1 /FL=gb:AB010710.1 gb:AF035776.1 gb:NM_002543.1	AF035776		NP_002534
210009_s_at	0.034317	gb:AF229796.1 /DEF=Homo sapiens golgi SNARE isoform B (GOSR2) mRNA, complete cds, alternatively spliced. /FEA=mRNA /GEN=GOSR2 /PROD=golgi SNARE isoform B /DB_XREF=gi:12711466 /UG=Hs.100651 golgi SNAP receptor complex member 2 /FL=gb:AF229796.1 gb:AF007548.1 gb:NM_004287.1	AF229796		NP_473363
210014_x_at	0.033381	gb:AF023266.1 /DEF=Homo sapiens NAD+-specific isocitrate dehydrogenase beta subunit isoform B mRNA, nuclear gene encoding mitochondrial product, complete cds. /FEA=mRNA /PROD=NAD+-specific isocitrate dehydrogenase betasubunit isoform B /DB_XREF=gi:4103447 /UG=Hs.155410 isocitrate dehydrogenase 3 (NAD+) beta /FL=gb:AF023266.1	AF023266		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210018_x_at	0.035611	gb:AB026118.1 /DEF=Homo sapiens mRNA for MALT1, complete cds. /FEA=mRNA /GEN=MALT1 /PROD=MALT1 /DB_XREF=gi:5706377 /UG=Hs.180566 mucosa associated lymphoid tissue lymphoma translocation gene 1 /FL=gb:AB026118.1	AB026118		NP_776216
210031_at	0.001054	gb:J04132.1 /DEF=Human T cell receptor zeta-chain mRNA, complete cds. /FEA=mRNA /PROD=T-cell receptor zeta chain /DB_XREF=gi:623041 /UG=Hs.97087 CD3Z antigen, zeta polypeptide (TiT3 complex) /FL=gb:NM_000734.1 gb:J04132.1	J04132		NP_000725
210044_s_at	0.004026	gb:BC002796.1 /DEF=Homo sapiens, lymphoblastic leukemia derived sequence 1, clone MGC:3943, mRNA, complete cds. /FEA=mRNA /PROD=lymphoblastic leukemia derived sequence 1 /DB_XREF=gi:12803898 /UG=Hs.46446 lymphoblastic leukemia derived sequence 1 /FL=gb:BC002796.1 gb:NM_005583.1	BC002796		NP_005574
210057_at	0.022708	gb:U32581.2 /DEF=Homo sapiens lambdaiota protein kinase C-interacting protein mRNA, complete cds. /FEA=mRNA /PROD=lambdaiota protein kinase C-interactingprotein /DB_XREF=gi:5542015 /UG=Hs.168052 KIAA0421 protein /FL=gb:U32581.2	U32581		NP_055907
210075_at	1.41E-04	gb:AF151074.1 /DEF=Homo sapiens HSPC240 mRNA, complete cds. /FEA=mRNA /PROD=HSPC240 /DB_XREF=gi:7106869 /UG=Hs.132744 hypothetical protein /FL=gb:AF151074.1 gb:NM_016496.1	AF151074		NP_057580

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210076_x_at	0.02145	gb:AF151813.1 /DEF=Homo sapiens CGI-55 protein mRNA, complete cds. /FEA=mRNA /PROD=CGI-55 protein /DB_XREF=gi:4929578 /UG=Hs.165998 PAI-1 mRNA-binding protein /FL=gb:AF151813.1	AF151813		NP_056455
210077_s_at	0.010061	gb:U30884.1 /DEF=Human splicing factor SRp40-2 (SRp40) mRNA, complete cds. /FEA=mRNA /GEN=SRp40 /PROD=SRp40-2 /DB_XREF=gi:1049089 /UG=Hs.166975 splicing factor, arginineserine-rich 5 /FL=gb:U30884.1	U30884		NP_008856
210088_x_at	2.17E-04	gb:M36172.1 /DEF=Human embryonic myosin alkali light chain (MLC1) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:188580 /UG=Hs.298161 myosin, light polypeptide 4, alkali; atrial, embryonic /FL=gb:M36172.1 gb:M24121.1 gb:NM_002476.1	M36172		NP_002467
210092_at	0.02145	gb:AF067173.1 /DEF=Homo sapiens Mago homolog mRNA, complete cds. /FEA=mRNA /PROD=Mago homolog /DB_XREF=gi:4894379 /UG=Hs.57904 mago-nashi (Drosophila) homolog, proliferation associated /FL=gb:AF035940.1 gb:AF067173.1 gb:NM_002370.2	AF067173		NP_002361
210097_s_at	0.033381	Consensus includes gb:AF130102.1 /DEF=Homo sapiens clone FLB2205 PRO0522 mRNA, complete cds. /FEA=mRNA /PROD=PRO0522 /DB_XREF=gi:11493507 /UG=Hs.106346 retinoic acid repressible protein /FL=gb:AF130102.1	AF130102		NP_057251

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210101_x_at	1.26E-04	gb:AF257318.1 /DEF=Homo sapiens SH3-containing protein SH3GLB1 mRNA, complete cds. /FEA=mRNA /PROD=SH3-containing protein SH3GLB1 /DB_XREF=gi:8896091 /UG=Hs.136309 SH3-containing protein SH3GLB1 /FL=gb:AF350371.1 gb:AF151819.1 gb:NM_016009.1 gb:AF257318.1	AF257318		NP_057093
210102_at	0.021045	gb:BC001234.1 /DEF=Homo sapiens, Similar to loss of heterozygosity, 11, chromosomal region 2, gene A, clone MGC:4904, mRNA, complete cds. /FEA=mRNA /PROD=Similar to loss of heterozygosity, 11,chromosomal region 2, gene A /DB_XREF=gi:12654782 /UG=Hs.152944 loss of heterozygosity, 11, chromosomal region 2, gene A /FL=gb:BC001234.1	BC001234		NP_055437
210113_s_at	0.003539	gb:AF310105.1 /DEF=Homo sapiens NALP1 mRNA, complete cds. /FEA=mRNA /PROD=NALP1 /DB_XREF=gi:11096302 /UG=Hs.104305 death effector filament-forming Ced-4-like apoptosis protein /FL=gb:AF310105.1 gb:AF229059.1 gb:AB023143.1 gb:NM_014922.1	AF310105		NP_127500
210117_at	0.00161	gb:AF311312.1 /DEF=Homo sapiens infertility-related sperm protein mRNA, complete cds. /FEA=mRNA /PROD=infertility-related sperm protein /DB_XREF=gi:10863767 /UG=Hs.153057 sperm associated antigen 1 /FL=gb:AF311312.1 gb:NM_003114.1	AF311312		NP_757367

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210119_at	3.26E-04	gb:U73191.1 /DEF=Human inward rectifier potassium channel (Kir1.3), complete cds. /FEA=mRNA /GEN=Kir1.3 /PROD=inward rectifier potassium channel /DB_XREF=gi:1765984 /UG=Hs.17287 potassium inwardly-rectifying channel, subfamily J, member 15 /FL=gb:U73191.1 gb:NM_002243.1	U73191		NP_733933
210128_s_at	0.003862	gb:U41070.1 /DEF=Human P2 purinergic receptor mRNA, complete cds. /FEA=mRNA /PROD=P2 purinergic receptor /DB_XREF=gi:1469913 /UG=Hs.28408 leukotriene b4 receptor (chemokine receptor-like 1) /FL=gb:U41070.1 gb:D89079.1	U41070		NP_000743
210131_x_at	3.26E-04	gb:D49737.1 /DEF=Homo sapiens mRNA for cytochrome b large subunit of complex II, complete cds. /FEA=mRNA /PROD=cytochrome b large subunit of complex II /DB_XREF=gi:2588778 /UG=Hs.3577 succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD /FL=gb:D49737.1	D49737		NP_002992
210136_at	5.03E-04	ESTs, Moderately similar to MBP_HUMAN Myelin basic protein (MBP) (Myelin A1 protein) (Myelin membrane encephalitogenic protein) [H.sapiens]	AW070431	Hs.409352	NP_002376
210140_at	0.00161	gb:AF031824.1 /DEF=Homo sapiens leukocystatin mRNA, complete cds. /FEA=mRNA /PROD=leukocystatin /DB_XREF=gi:3252857 /UG=Hs.143212 cystatin F (leukocystatin) /FL=gb:AF031824.1 gb:AF036342.1 gb:AB015225.1 gb:NM_003650.1	AF031824		NP_003641

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210142_x_at	2.05E-04	gb:AF117234.1 /DEF=Homo sapiens flotillin mRNA, complete cds. /FEA=mRNA /PROD=flotillin /DB_XREF=gi:6563241 /UG=Hs.179986 flotillin 1 /FL=gb:AF117234.1	AF117234		NP_005794
210144_at	0.021165	Consensus includes gb:AK000851.1 /DEF=Homo sapiens cDNA FLJ20844 fis, clone ADKA01904. /FEA=mRNA /DB_XREF=gi:7021173 /UG=Hs.20017 chromosome 22 open reading frame 4 /FL=gb:BC002743.1	AK000851		NP_055161
210145_at	0.00161	gb:M68874.1 /DEF=Homo sapiens phosphatidylcholine 2-acylhydrolase (cPLA2) mRNA, complete cds. /FEA=mRNA /GEN=cPLA2 /PROD=phosphatidylcholine 2-acylhydrolase /DB_XREF=gi:190003. /UG=Hs.211587 phospholipase A2, group IVA (cytosolic, calcium-dependent) /FL=gb:M68874.1 gb:M72393.1	M68874		NP_077734
210146_x_at	0.016934	gb:AF004231.1 /DEF=Homo sapiens monocytemacrophage Ig-related receptor MIR-10 (MIR cl-10) mRNA, complete cds. /FEA=mRNA /GEN=MIR cl-10 /PROD=MIR-10 /DB_XREF=gi:2343110 /UG=Hs.22405 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2 /FL=gb:AF000574.1 gb:AF004231.1 gb:AF011565.1	AF004231		NP_005865
210149_s_at	1.26E-04	gb:AF061735.1 /DEF=Homo sapiens clone 023e08 My032 protein mRNA, complete cds. /FEA=mRNA /PROD=My032 protein /DB_XREF=gi:12002005 /UG=Hs.307099 Homo sapiens clone 023e08 My032 protein mRNA, complete cds /FL=gb:AF061735.1	AF061735		NP_006347

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210153_s_at	2.05E-04	gb:M55905.1 /DEF=Human mitochondrial NAD(P)+ dependent malic enzyme mRNA, complete cds. /FEA=mRNA /GEN=NAD(P)+ -dependent malic enzyme /PROD=mitochondrial NAD(P)+ -dependent malic enzyme /DB_XREF=gi:187299 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:NM_002396.2 gb:M55905.1	M55905		NP_002387
210154_at	5.64E-04	gb:M55905.1 /DEF=Human mitochondrial NAD(P)+ dependent malic enzyme mRNA, complete cds. /FEA=mRNA /GEN=NAD(P)+ -dependent malic enzyme /PROD=mitochondrial NAD(P)+ -dependent malic enzyme /DB_XREF=gi:187299 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:NM_002396.2 gb:M55905.1	M55905		NP_002387
210156_s_at	1.26E-04	gb:D25547.1 /DEF=Homo sapiens mRNA for PIMT isozyme I, complete cds. /FEA=mRNA /PROD=PIMT isozyme I /DB_XREF=gi:1332402 /UG=Hs.79137 protein-L-isoaspartate (D-aspartate) O-methyltransferase /FL=gb:D25547.1	D25547		NP_005380
210162_s_at	0.041055	gb:U08015.1 /DEF=Human NF-ATc mRNA, complete cds. /FEA=mRNA /PROD=NF-ATc /DB_XREF=gi:500631 /UG=Hs.96149 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 /FL=gb:U08015.1	U08015		NP_765978

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210176_at	0.003171	gb:AL050262.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I0682 (from clone DKFZp564I0682); complete cds. /FEA=mRNA /GEN=DKFZp564I0682 /PROD=hypothetical protein /DB_XREF=gi:4886482 /UG=Hs.2474 toll-like receptor 1 /FL=gb:U88540.1 gb:D13637.1 gb:NM_003263.1 gb:AL050262.1	AL050262		NP_003254
210187_at	0.02145	gb:BC005147.1 /DEF=Homo sapiens, FK506-binding protein 1A (12kD), clone MGC:2167, mRNA, complete cds. /FEA=mRNA /PROD=FK506-binding protein 1A (12kD) /DB_XREF=gi:13477342 /UG=Hs.752 FK506-binding protein 1A (12kD) /FL=gb:BC005147.1	BC005147		NP_463460
210191_s_at	1.26E-04	gb:BC002447.1 /DEF=Homo sapiens, Similar to putative homeodomain transcription factor, clone MGC:1307, mRNA, complete cds. /FEA=mRNA /PROD=Similar to putative homeodomain transcriptionfactor /DB_XREF=gi:12803264 /UG=Hs.123637 putative homeodomain transcription factor /FL=gb:BC002447.1	BC002447		NP_006599
210200_at	7.58E-04	gb:BC000108.1 /DEF=Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Nedd-4-like ubiquitin-protein ligase /DB_XREF=gi:12652716 /UG=Hs.19301 Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds /FL=gb:BC000108.1	BC000108		NP_008945

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210205_at	0.010205	gb:AB026730.1 /DEF=Homo sapiens B3GALT4 mRNA for beta-1,3-galactosyltransferase-4, complete cds. /FEA=mRNA /GEN=B3GALT4 /PROD=beta-1,3-galactosyltransferase-4 /DB_XREF=gi:6683013 /UG=Hs.21495 UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4 /FL=gb:AF063595.1 gb:NM_003782.1 gb:AB026730.1	AB026730		NP_003773
210213_s_at	5.03E-04	gb:AF022229.1 /DEF=Homo sapiens translation initiation factor 6 (eIF6) mRNA, complete cds. /FEA=mRNA /GEN=eIF6 /PROD=translation initiation factor 6 /DB_XREF=gi:2809382 /UG=Hs.5215 integrin beta 4 binding protein /FL=gb:AF022229.1	AF022229		NP_002203
210218_s_at	0.012818	gb:U36501.1 /DEF=Human SP100 B (SP100-B) mRNA, complete cds. /FEA=mRNA /GEN=SP100-B /PROD=SP100-B /DB_XREF=gi:1173655 /UG=Hs.77617 nuclear antigen Sp100 /FL=gb:U36501.1	U36501		NP_003104
210225_x_at	0.013223	gb:AF009635.1 /DEF=Homo sapiens clone 17.7 immunoglobulin-like transcript 5 protein mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 5 protein /DB_XREF=gi:2662429 /UG=Hs.105928 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 /FL=gb:AF000575.1 gb:U91928.1 gb:AF025533.1 gb:AF009632.1 gb:AF009633.1 gb:AF009635.1 gb:AF009638.1 gb:AF009639.1 gb:AF009641.1 gb:AF009642.1 gb:AF031553.1 gb:AF031554.1 gb:AF031555.1 gb:AF031556.1 gb:NM_006864.1	AF009635		NP_006855

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210231_x_at	1.26E-04	gb:D45198.1 /DEF=Human mRNA for template acyivating factor-I alpha, complete cds. /FEA=mRNA /GEN=set /PROD=template acyivating factor-I alpha /DB_XREF=gi:971271 /UG=Hs.145279 SET translocation (myeloid leukemia-associated) /FL=gb:D45198.1	D45198		NP_003002
210240_s_at	0.003529	gb:U20498.1 /DEF=Human p19 protein mRNA, complete cds. /FEA=mRNA /PROD=p19 protein /DB_XREF=gi:1161921 /UG=Hs.29656 cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4) /FL=gb:U40343.1 gb:U20498.1 gb:BC001822.1 gb:U49399.1 gb:NM_001800.1	U20498		NP_524145
210241_s_at	0.027239	gb:AB007458.1 /DEF=Homo sapiens mRNA for P53TG1-D, complete cds. /FEA=mRNA /GEN=P53TG1 /PROD=P53TG1-D /DB_XREF=gi:5006272 /UG=Hs.306301 Homo sapiens mRNA for P53TG1-D, complete cds /FL=gb:AB007458.1	AB007458		NP_009164
210244_at	1.41E-04	gb:U19970.1 /DEF=Human antimicrobial LPS-binding protein CAP18 precursor mRNA, complete cds. /FEA=mRNA /PROD=CAP18 precursor /DB_XREF=gi:643476 /UG=Hs.51120 cathelicidin antimicrobial peptide /FL=gb:NM_004345.1 gb:U19970.1	U19970		NP_004336
210249_s_at	0.00228	gb:U59302.1 /DEF=Human steroid receptor coactivator-1 F-SRC-1 mRNA, complete cds. /FEA=mRNA /PROD=steroid receptor coactivator-1 F-SRC-1 /DB_XREF=gi:1480645 /UG=Hs.74002 nuclear receptor coactivator 1 /FL=gb:U40396.1 gb:U59302.1 gb:U90661.1 gb:NM_003743.1	U59302		NP_671766

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210254_at	1.41E-04	gb:L35848.1 /DEF=Homo sapiens IgE receptor beta chain (HTm4) mRNA, complete cds. /FEA=mRNA /GEN=HTm4 /PROD=IgE receptor beta subunit /DB_XREF=gi:561638 /UG=Hs.99960 membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific) /FL=gb:NM_006138.1 gb:L35848.1	L35848		NP_006129
210255_at	0.019999	gb:U84138.1 /DEF=Homo sapiens DNA repair protein RAD51B mRNA, complete cds. /FEA=mRNA /PROD=DNA repair protein RAD51B /DB_XREF=gi:2801404 /UG=Hs.100669 RAD51 (S. cerevisiae)-like 1 /FL=gb:NM_002877.1 gb:U92074.1 gb:U84138.1	U84138		NP_598194
210260_s_at	0.02145	gb:BC005352.1 /DEF=Homo sapiens, TNF-induced protein, clone MGC:12451, mRNA, complete cds. /FEA=mRNA /PROD=TNF-induced protein /DB_XREF=gi:13529163 /UG=Hs.17839 TNF-induced protein /FL=gb:BC005352.1 gb:AF099935.1	BC005352		NP_055165
210275_s_at	2.05E-04	gb:AF062347.1 /DEF=Homo sapiens zinc finger protein 216 splice variant 2 (ZNF216) mRNA, complete cds. /FEA=mRNA /GEN=ZNF216 /PROD=zinc finger protein 216 splice variant 2 /DB_XREF=gi:3643810 /UG=Hs.3776 zinc finger protein 216 /FL=gb:AF062347.1	AF062347		NP_005998
210279_at	0.033381	gb:AF261135.1 /DEF=Homo sapiens GPR18-iso mRNA, complete cds. /FEA=mRNA /PROD=GPR18-iso /DB_XREF=gi:12005919 /UG=Hs.88269 Homo sapiens clone IMAGE:1837189, mRNA sequence /FL=gb:AF261135.1	AF261135		NP_005283

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210283_x_at	7.58E-04	gb:BC005295.1 /DEF=Homo sapiens, Similar to polyadenylate binding protein-interacting protein 1, clone MGC:12360, mRNA, complete cds. /FEA=mRNA /PROD=Similar to polyadenylate bindingprotein-interacting protein 1 /DB_XREF=gi:13529010 /UG=Hs.109643 polyadenylate binding protein-interacting protein 1 /FL=gb:BC005295.1	BC005295		NP_006442
210284_s_at	0.041055	gb:AF241230.1 /DEF=Homo sapiens TAK1-binding protein 2 mRNA, complete cds. /FEA=mRNA /PROD=TAK1-binding protein 2 /DB_XREF=gi:7677465 /UG=Hs.109727 TAK1-binding protein 2; KIAA0733 protein /FL=gb:AF241230.1	AF241230		NP_663317
210285_x_at	0.013223	gb:BC000383.1 /DEF=Homo sapiens, Wilms tumour 1-associating protein, clone MGC:8419, mRNA, complete cds. /FEA=mRNA /PROD=Wilms tumour 1-associating protein /DB_XREF=gi:12653228 /UG=Hs.119 Wilms tumour 1-associating protein /FL=gb:BC000383.1 gb:BC004432.1	BC000383		NP_690597
210288_at	0.00161	gb:AF081675.1 /DEF=Homo sapiens ITIM-containing receptor MAFA-L mRNA, complete cds. /FEA=mRNA /PROD=ITIM-containing receptor MAFA-L /DB_XREF=gi:3421400 /UG=Hs.146322 killer cell lectin-like receptor subfamily G, member 1 /FL=gb:AF034952.1 gb:AF081675.1 gb:AF097358.1 gb:NM_005810.1	AF081675		NP_005801

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210293_s_at	1.26E-04	gb:BC005032.1 /DEF=Homo sapiens, Sec23 (S. cerevisiae) homolog B, clone MGC:12666, mRNA, complete cds. /FEA=mRNA /PROD=Sec23 (S. cerevisiae) homolog B /DB_XREF=gi:13477148 /UG=Hs.173497 Sec23 (S. cerevisiae) homolog B /FL=gb:BC005032.1	BC005032		NP_116781
210296_s_at	5.03E-04	gb:BC005375.1 /DEF=Homo sapiens, peroxisomal membrane protein 3 (35kD, Zellweger syndrome), clone MGC:12491, mRNA, complete cds. /FEA=mRNA /PROD=peroxisomal membrane protein 3 (35kD, Zellwegersyndrome) /DB_XREF=gi:13529226 /UG=Hs.180612 peroxisomal membrane protein 3 (35kD, Zellweger syndrome) /FL=gb:BC005375.1	BC005375		NP_000309
210299_s_at	7.57E-04	gb:AF063002.1 /DEF=Homo sapiens LIM protein SLIMMER mRNA, complete cds. /FEA=mRNA /PROD=LIM protein SLIMMER /DB_XREF=gi:3859848 /UG=Hs.239069 four and a half LIM domains 1 /FL=gb:AF098518.1 gb:AF063002.1	AF063002		NP_001440
210312_s_at	1.26E-04	gb:BC002640.1 /DEF=Homo sapiens, Similar to uterine protein, clone MGC:4279, mRNA, complete cds. /FEA=mRNA /PROD=Similar to uterine protein /DB_XREF=gi:12803610 /UG=Hs.4187 hypothetical protein 24636 /FL=gb:BC002640.1	BC002640		NP_777547

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210314_x_at	2.05E-04	gb:AF114013.1 /DEF=Homo sapiens tumor necrosis factor-related death ligand-1gamma mRNA, complete cds. /FEA=mRNA /PROD=tumor necrosis factor-related deathligand 1gamma /DB_XREF=gi:7328557 /UG=Hs.54673 tumor necrosis factor (ligand) superfamily, member 13 /FL=gb:AF136294.1 gb:AF114013.1	AF114013		NP_742086
210340_s_at	0.001116	gb:BC002635.1 /DEF=Homo sapiens, colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage), clone MGC:3848, mRNA, complete cds. /FEA=mRNA /PROD=colony stimulating factor 2 receptor, alpha,low-affinity (granulocyte-macrophage) /DB_XREF=gi:12803600 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:BC002635.1	BC002635		NP_758452
210356_x_at	0.003539	gb:BC002807.1 /DEF=Homo sapiens, membrane-spanning 4-domains, subfamily A, member 2, clone MGC:3969, mRNA, complete cds. /FEA=mRNA /PROD=membrane-spanning 4-domains, subfamily A, member2 /DB_XREF=gi:12803920 /UG=Hs.89751 membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide) /FL=gb:NM_021950.1 gb:BC002807.1	BC002807		NP_690606

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210360_s_at	0.037567	gb:AF116674.1 /DEF=Homo sapiens PRO1941 mRNA, complete cds. /FEA=mRNA /PROD=PRO1941 /DB_XREF=gi:7959846 /UG=Hs.77694 KIAA0429 gene product /FL=gb:AF116674.1	AF116674		NP_055566
210367_s_at	0.02145	gb:AF010316.1 /DEF=Homo sapiens Pig12 (PIG12) mRNA, complete cds. /FEA=mRNA /GEN=PIG12 /PROD=Pig12 /DB_XREF=gi:2415307 /UG=Hs.146688 prostaglandin E synthase /FL=gb:AF010316.1	AF010316		NP_004869
210386_s_at	7.58E-04	gb:BC001906.1 /DEF=Homo sapiens, Similar to metaxin 1, clone MGC:2518, mRNA, complete cds. /FEA=mRNA /PROD=Similar to metaxin 1 /DB_XREF=gi:12804906 /UG=Hs.247551 metaxin 1 /FL=gb:BC001906.1	BC001906		NP_002446
210395_x_at	3.26E-04	gb:AF116676.1 /DEF=Homo sapiens PRO1957 mRNA, complete cds. /FEA=mRNA /PROD=PRO1957 /DB_XREF=gi:7959850 /UG=Hs.298161 myosin, light polypeptide 4, alkali; atrial, embryonic /FL=gb:AF116676.1	AF116676		NP_002467
210401_at	0.005852	gb:U45448.1 /DEF=Human P2x1 receptor mRNA, complete cds. /FEA=mRNA /PROD=P2X1 receptor /DB_XREF=gi:1314356 /UG=Hs.41735 purinergic receptor P2X, ligand-gated ion channel, 1 /FL=gb:U45448.1 gb:AF020498.1 gb:NM_002558.1	U45448		NP_002549
210406_s_at	1.26E-04	gb:AL136727.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566K144 (from clone DKFZp566K144); complete cds. /FEA=mRNA /GEN=DKFZp566K144 /PROD=hypothetical protein /DB_XREF=gi:12052972 /UG=Hs.5636 RAB6A, member RAS oncogene family /FL=gb:AL136727.1	AL136727		NP_115520

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210407_at	0.013088	gb:AF070670.1 /DEF=Homo sapiens protein phosphatase 2C alpha 2 mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 2C alpha 2 /DB_XREF=gi:3372871 /UG=Hs.57764 protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform /FL=gb:AF070670.1	AF070670		NP_808821
210422_x_at	0.02145	gb:D50402.1 /DEF=Human mRNA for NRAMP1, complete cds. /FEA=mRNA /PROD=Nramp /DB_XREF=gi:1000996 /UG=Hs.182611 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1 /FL=gb:D50402.1 gb:D50403.1 gb:NM_000578.1 gb:L32185.1	D50402		NP_000569
210434_x_at	0.010205	gb:AF151056.1 /DEF=Homo sapiens HSPC222 mRNA, complete cds. /FEA=mRNA /PROD=HSPC222 /DB_XREF=gi:7106833 /UG=Hs.323093 Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRNA, complete cds /FL=gb:AF151056.1	AF151056		NP_006685
210438_x_at	0.016934	gb:M25077.1 /DEF=Human SS-ARo ribonucleoprotein autoantigen 60 kd subunit mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:387656 /UG=Hs.554 Sjogren syndrome antigen A2 (60kD, ribonucleoprotein autoantigen SS-ARo) /FL=gb:M25077.1	M25077		NP_004591

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210453_x_at	5.03E-04	gb:AL050277.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566G013 (from clone DKFZp566G013); complete cds. /FEA=mRNA /GEN=DKFZp566G013 /PROD=hypothetical protein /DB_XREF=gi:4886454 /UG=Hs.107476 ATP synthase, H ⁺ transporting, mitochondrial F1F0, subunit g /FL=gb:AL050277.1	AL050277		NP_006467
210460_s_at	1.26E-04	gb:AB033605.1 /DEF=Homo sapiens mRNA for pUb-R5, complete cds. /FEA=mRNA /GEN=hpUb-R5 /PROD=pUb-R5 /DB_XREF=gi:8918352 /UG=Hs.148495 proteasome (prosome, macropain) 26S subunit, non-ATPase, 4 /FL=gb:AB033605.1	AB033605		NP_722544
210479_s_at	0.003539	gb:L14611.1 /DEF=Human transcription factor RZR-alpha mRNA, complete cds. /FEA=mRNA /PROD=transcription factor /DB_XREF=gi:348240 /UG=Hs.2156 RAR-related orphan receptor A /FL=gb:L14611.1 gb:NM_002943.1 gb:U04899.1	L14611		NP_599024

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210483_at	0.04734	gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, complete cds. /FEA=mRNA /PROD=Similar to tumor necrosis factor receptorsuperfamily, member 10c, decoy without an intracellulardomain /DB_XREF=gi:13477162 /UG=Hs.253829 Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, complete cds /FL=gb:BC005043.1	BC005043		
210499_s_at	0.026891	gb:AB041834.1 /DEF=Homo sapiens mRNA for PQBP-1bc, complete cds. /FEA=mRNA /PROD=PQBP-1bc /DB_XREF=gi:10801589 /UG=Hs.30570 polyglutamine binding protein 1 /FL=gb:AB041834.1	AB041834		NP_005701
210504_at	0.001477	gb:U65404.1 /DEF=Human erythroid-specific transcription factor EKLF mRNA, complete cds. /FEA=mRNA /PROD=erythroid-specific transcription factor EKLF /DB_XREF=gi:1552995 /UG=Hs.37860 Kruppel-like factor 1 (erythroid) /FL=gb:NM_006563.1 gb:U65404.1	U65404		NP_006554
210506_at	0.003741	gb:U11282.1 /DEF=Human alpha(1,3)fucosyltransferase mRNA, complete cds. /FEA=mRNA /PROD=alpha(1,3)fucosyltransferase /DB_XREF=gi:520524 /UG=Hs.457 fucosyltransferase 7 (alpha (1,3) fucosyltransferase) /FL=gb:NM_004479.1 gb:U08112.1 gb:U11282.1	U11282		NP_004470

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210512_s_at	0.003862	gb:AF022375.1 /DEF=Homo sapiens vascular endothelial growth factor mRNA, complete cds. /FEA=mRNA /PROD=vascular endothelial growth factor /DB_XREF=gi:3719220 /UG=Hs.73793 vascular endothelial growth factor /FL=gb:M32977.1 gb:AF022375.1 gb:NM_003376.1 gb:AB021221.1 gb:AF091352.1	AF022375		NP_003367
210532_s_at	1.26E-04	gb:AF116639.1 /DEF=Homo sapiens PRO1574 mRNA, complete cds. /FEA=mRNA /PROD=PRO1574 /DB_XREF=gi:7959779 /UG=Hs.109052 chromosome 14 open reading frame 2 /FL=gb:AF116639.1	AF116639		
210542_s_at	0.004341	gb:BC000585.1 /DEF=Homo sapiens, Similar to solute carrier family 21 (organic anion transporter), member 11, clone MGC:659, mRNA, complete cds. /FEA=mRNA /PROD=Similar to solute carrier family 21 (organicanion transporter), member 11 /DB_XREF=gi:12653614 /UG=Hs.14805 solute carrier family 21 (organic anion transporter), member 11 /FL=gb:BC000585.1	BC000585		NP_037404
210556_at	0.0174	gb:U85430.1 /DEF=Human transcription factor NFATx4 mRNA, complete cds. /FEA=mRNA /PROD=transcription factor NFATx4 /DB_XREF=gi:1835590 /UG=Hs.172674 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 /FL=gb:U85430.1	U85430		NP_775188

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210573_s_at	0.042536	gb:BC004424.1 /DEF=Homo sapiens, clone MGC:3538, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3538) /DB_XREF=gi:13325209 /UG=Hs.250745 polymerase (RNA) III (DNA directed) (62kD) /FL=gb:BC004424.1	BC004424		NP_006459
210582_s_at	0.016934	gb:AL117466.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586K0922 (from clone DKFZp586K0922); complete cds. /FEA=mRNA /GEN=DKFZp586K0922 /PROD=hypothetical protein /DB_XREF=gi:5911930 /UG=Hs.278027 LIM domain kinase 2 /FL=gb:AL117466.1 gb:NM_016733.1	AL117466		NP_057952
210592_s_at	0.033381	gb:M55580.1 /DEF=Human spermidinespermine N1-acetyltransferase mRNA, complete cds. /FEA=mRNA /GEN=spermidinespermine N1-acetyltransferase /PROD=spermidinespermine N1-acetyltransferase /DB_XREF=gi:338335 /UG=Hs.28491 spermidinespermine N1-acetyltransferase /FL=gb:M55580.1	M55580		
210598_at	0.004341	gb:AF130051.1 /DEF=Homo sapiens clone FLB3535 PRO0898 mRNA, complete cds. /FEA=mRNA /PROD=PRO0898 /DB_XREF=gi:11493408 /UG=Hs.306960 Homo sapiens clone FLB3535 PRO0898 mRNA, complete cds /FL=gb:AF130051.1	AF130051		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210608_s_at	0.017832	gb:BC001899.1 /DEF=Homo sapiens, Similar to fucosyltransferase 2 (secretor status included), clone MGC:2062, mRNA, complete cds. /FEA=mRNA /PROD=Similar to fucosyltransferase 2 (secretor status included) /DB_XREF=gi:12804894 /UG=Hs.46328 fucosyltransferase 2 (secretor status included) /FL=gb:BC001899.1	BC001899		NP_000502
210621_s_at	0.003171	gb:M23612.1 /DEF=Human GTPase-activating protein (GAP) mRNA, complete cds. /FEA=mRNA /GEN=GAP /PROD=GTPase-activating protein /DB_XREF=gi:182973 /UG=Hs.758 RAS p21 protein activator (GTPase activating protein) 1 /FL=gb:NM_022650.1 gb:M23612.1	M23612		NP_072179
210624_s_at	0.010205	gb:BC000109.1 /DEF=Homo sapiens, Similar to ilvB (bacterial acetolactate synthase)-like, clone MGC:1269, mRNA, complete cds. /FEA=mRNA /PROD=Similar to ilvB (bacterial acetolactatesynthase)-like /DB_XREF=gi:12652718 /UG=Hs.78880 ilvB (bacterial acetolactate synthase)-like /FL=gb:BC000109.1	BC000109		NP_789796
210629_x_at	0.004341	gb:AF000425.1 /DEF=Homo sapiens LST1 mRNA, cLST1A splice variant, complete cds. /FEA=mRNA /GEN=LST1 /DB_XREF=gi:2145065 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:AF000425.1	AF000425		NP_009092
210638_s_at	1.26E-04	gb:AF176704.1 /DEF=Homo sapiens F-box protein FBX9 mRNA, complete cds. /FEA=mRNA /PROD=F-box protein FBX9 /DB_XREF=gi:6103646 /UG=Hs.11050 F-box only protein 9 /FL=gb:AF176704.1	AF176704		NP_258442

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210639_s_at	2.05E-04	gb:AF293841.1 /DEF=Homo sapiens apoptosis-related protein (APG5L) mRNA, complete cds, alternatively spliced. /FEA=mRNA /GEN=APG5L /PROD=apoptosis-related protein /DB_XREF=gi:12006863 /UG=Hs.11171 APG5 (autophagy 5, <i>S. cerevisiae</i>)-like /FL=gb:AF293841.1	AF293841		NP_004840
210644_s_at	7.58E-04	gb:AF109683.1 /DEF=Homo sapiens leukocyte-associated Ig-like receptor 1b mRNA, complete cds. /FEA=mRNA /PROD=leukocyte-associated Ig-like receptor 1b /DB_XREF=gi:6563041 /UG=Hs.115808 leukocyte-associated Ig-like receptor 1 /FL=gb:AF251509.2 gb:NM_021706.1 gb:AF109683.1	AF109683		NP_068354
210646_x_at	0.007779	gb:BC001675.1 /DEF=Homo sapiens, ribosomal protein L13a, clone MGC:2546, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L13a /DB_XREF=gi:12804528 /UG=Hs.119122 ribosomal protein L13a /FL=gb:BC001675.1 gb:BC001836.1	BC001675		NP_036555
210647_x_at	0.004341	gb:AF102988.1 /DEF=Homo sapiens Ca ²⁺ -independent phospholipase A2 short isoform (iPLA2) mRNA, complete cds. /FEA=mRNA /GEN=iPLA2 /PROD=Ca ²⁺ -independent phospholipase A2 short isoform /DB_XREF=gi:5305593 /UG=Hs.120360 phospholipase A2, group VI (cytosolic, calcium-independent) /FL=gb:AF102988.1	AF102988		NP_003551

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210648_x_at	1.26E-04	gb:AB047360.1 /DEF=Homo sapiens SNX 3A mRNA for sorting nexin 3A, complete cds. /FEA=mRNA /GEN=SNX 3A /PROD=sorting nexin 3A /DB_XREF=gi:12957159 /UG=Hs.12102 sorting nexin 3 /FL=gb:AB047360.1	AB047360		NP_690041
210667_s_at	6.93E-04	gb:D86062.1 /DEF=Human mRNA for KNP-Ib, complete cds. /FEA=mRNA /GEN=KNP-I /PROD=KNP-Ib /DB_XREF=gi:1545814 /UG=Hs.182423 ES1 (zebrafish) protein, human homolog of /FL=gb:D86062.1	D86062		NP_004640
210679_x_at	0.003171	gb:BC002629.1 /DEF=Homo sapiens, clone MGC:3878, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3878) /DB_XREF=gi:12803588 /UG=Hs.211563 B-cell CLL lymphoma 7A /FL=gb:BC002629.1	BC002629		
210685_s_at	0.004026	gb:AB028839.1 /DEF=Homo sapiens HDNB1(homzygously deleted in neuroblastoma-1)UFD2 mRNA, complete cds. /FEA=mRNA /GEN=HDNB1UFD2 /PROD=homzygously deleted in neuroblastoma-1UFD2 /DB_XREF=gi:13516466 /UG=Hs.24594 ubiquitination factor E4B (homologous to yeast UFD2) /FL=gb:AB028839.1	AB028839		NP_006039
210686_x_at	1.26E-04	gb:BC001407.1 /DEF=Homo sapiens, Similar to cytochrome c-like antigen, clone MGC:2960, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cytochrome c-like antigen /DB_XREF=gi:12655110 /UG=Hs.253070 Homo sapiens, Similar to cytochrome c-like antigen, clone MGC:2960, mRNA, complete cds /FL=gb:BC001407.1	BC001407		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210691_s_at	5.03E-04	gb:AF275803.1 /DEF=Homo sapiens PNAS-107 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-107 /DB_XREF=gi:10834769 /UG=Hs.27258 calcyclin binding protein /FL=gb:AF275803.1	AF275803		NP_055227
210692_s_at	0.001483	gb:BC003163.1 /DEF=Homo sapiens, Similar to selectively expressed in embryonic epithelia protein-1, clone MGC:4220, mRNA, complete cds. /FEA=mRNA /PROD=Similar to selectively expressed in embryonicepithelia protein-1 /DB_XREF=gi:13111980 /UG=Hs.274453 hypothetical protein DKFZp762A227 /FL=gb:BC003163.1	BC003163		
210706_s_at	6.68E-04	gb:BC000213.1 /DEF=Homo sapiens, ring finger protein 24, clone MGC:1815, mRNA, complete cds. /FEA=mRNA /PROD=ring finger protein 24 /DB_XREF=gi:12802985 /UG=Hs.30524 ring finger protein 24 /FL=gb:BC000213.1	BC000213		NP_009150
210715_s_at	0.016934	gb:AF027205.1 /DEF=Homo sapiens Kunitz-type protease inhibitor (kop) mRNA, complete cds. /FEA=mRNA /GEN=kop /PROD=Kunitz-type protease inhibitor /DB_XREF=gi:2598967 /UG=Hs.31439 serine protease inhibitor, Kunitz type, 2 /FL=gb:AF027205.1	AF027205		NP_066925
210716_s_at	1.26E-04	gb:M97501.1 /DEF=Human cytoplasmic linker protein-170 alpha-2 (CLIP-170) mRNA, complete cds. /FEA=mRNA /PROD=cytoplasmic linker protein-170 alpha-2 /DB_XREF=gi:180621 /UG=Hs.31638 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) /FL=gb:M97501.1	M97501		NP_002947

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210718_s_at	0.037567	gb:AF119889.1 /DEF=Homo sapiens PRO2667 mRNA, complete cds. /FEA=mRNA /PROD=PRO2667 /DB_XREF=gi:7770214 /UG=Hs.321170 Homo sapiens PRO2667 mRNA, complete cds /FL=gb:AF119889.1	AF119889		NP_005812
210756_s_at	0.005768	gb:AF308601.1 /DEF=Homo sapiens NOTCH 2 (N2) mRNA, complete cds. /FEA=mRNA /GEN=N2 /PROD=NOTCH 2 /DB_XREF=gi:11275977 /UG=Hs.8121 Notch (Drosophila) homolog 2 /FL=gb:AF308601.1	AF308601		
210759_s_at	0.033381	gb:M64992.1 /DEF=Human prosomal protein. P30-33K (pros-30) mRNA, complete cds. /FEA=mRNA /GEN=pros-30 /PROD=prosomal protein P30-33K /DB_XREF=gi:190446 /UG=Hs.82159 proteasome (prosome, macropain) subunit, alpha type, 1 /FL=gb:M64992.1	M64992		NP_683877
210772_at	0.013223	gb:M88107.1 /DEF=Human formyl peptide receptor (FPR2) mRNA, complete cds. /FEA=mRNA /GEN=FPR2 /PROD=formyl peptide receptor /DB_XREF=gi:189862 /UG=Hs.99855 formyl peptide receptor-like 1 /FL=gb:M76672.1 gb:M84562.1 gb:M88107.1 gb:U81501.1 gb:AF054013.1 gb:NM_001462.1	M88107		NP_001453
210775_x_at	6.79E-04	gb:AB015653.1 /DEF=Homo sapiens mRNA for caspase-9 beta, complete cds. /FEA=mRNA /PROD=caspase-9 beta /DB_XREF=gi:5103139 /UG=Hs.100641 caspase 9, apoptosis-related cysteine protease /FL=gb:AF093130.1 gb:AF110376.1 gb:AB015653.1 gb:AB020979.1	AB015653		NP_127463

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210788_s_at	1.26E-04	gb:AF126782.1 /DEF=Homo sapiens retinal short-chain dehydrogenasereductase retSDR4 mRNA, complete cds. /FEA=mRNA /PROD=retinal short-chain dehydrogenasereductaseretSDR4 /DB_XREF=gi:6318547 /UG=Hs.109201 CGI-86 protein /FL=gb:AF126782.1	AF126782		
210789_x_at	0.042536	gb:L00692.1 /DEF=Human carcinoembryonic antigen (CGM1) mRNA, complete cds. /FEA=mRNA /GEN=CGM1 /PROD=carcinoembryonic antigen /DB_XREF=gi:180226 /UG=Hs.11 carcinoembryonic antigen-related cell adhesion molecule 3 /FL=gb:L00692.1	L00692		NP_001806
210793_s_at	5.03E-04	gb:U41815.1 /DEF=Human nucleoporin 98 (NUP98) mRNA, complete cds. /FEA=mRNA /GEN=NUP98 /PROD=nucleoporin 98 /DB_XREF=gi:1184172 /UG=Hs.112255 nucleoporin 98kD /FL=gb:U41815.1	U41815		NP_624358
210835_s_at	0.016934	gb:AF222711.1 /DEF=Homo sapiens ribeye mRNA, complete cds. /FEA=mRNA /PROD=ribeye /DB_XREF=gi:12034652 /UG=Hs.171391 C-terminal binding protein 2 /FL=gb:AF222711.1 gb:NM_022802.1	AF222711		NP_073713
210840_s_at	0.007779	gb:D29640.1 /DEF=Human mRNA for KIAA0051 gene, complete cds. /FEA=mRNA /GEN=KIAA0051 /DB_XREF=gi:473930 /UG=Hs.1742 IQ motif containing GTPase activating protein 1 /FL=gb:D29640.1	D29640		NP_003861

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210844_x_at	5.03E-04	gb:D14705.1 /DEF=Human mRNA for alpha-catenin, complete cds. /FEA=mRNA /PROD=human alpha-catenin /DB_XREF=gi:415305 /UG=Hs.178452 catenin (cadherin-associated protein), alpha 1 (102kD) /FL=gb:BC000385.1 gb:D14705.1 gb:D13866.1	D14705		NP_001894
210845_s_at	0.00161	gb:U08839.1 /DEF=Human urokinase-type plasminogen activator receptor mRNA, complete cds. /FEA=mRNA /PROD=urokinase-type plasminogen activator receptor /DB_XREF=gi:517197 /UG=Hs.179657 plasminogen activator, urokinase receptor /FL=gb:U08839.1	U08839		NP_002650
210858_x_at	0.013223	gb:U26455.1 /DEF=Human phosphatidylinositol 3-kinase homolog (ATM) mRNA, complete cds. /FEA=mRNA /GEN=ATM /PROD=phosphatidylinositol 3-kinase homolog /DB_XREF=gi:870785 /UG=Hs.194382 ataxia telangiectasia mutated (includes complementation groups A, C and D) /FL=gb:U26455.1	U26455		NP_612150
210859_x_at	0.001618	gb:AF077973.1 /DEF=Homo sapiens clone 36 CLN3 protein (CLN3) mRNA, complete cds. /FEA=mRNA /GEN=CLN3 /PROD=CLN3 protein /DB_XREF=gi:5801864 /UG=Hs.194660 ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease) /FL=gb:AF077973.1	AF077973		NP_000077

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210872_x_at	0.022752	gb:BC001152.1 /DEF=Homo sapiens, growth arrest-specific 7, clone MGC:1347, mRNA, complete cds. /FEA=mRNA /PROD=growth arrest-specific 7 /DB_XREF=gi:12654630 /UG=Hs.226133 growth arrest-specific 7 /FL=gb:BC001152.1	BC001152		NP_005881
210875_s_at	0.001848	gb:U12170.1 /DEF=Human zinc finger homeodomain protein mRNA, complete cds. /FEA=mRNA /PROD=zinc finger homeodomain protein /DB_XREF=gi:529172 /UG=Hs.232068 transcription factor 8 (represses interleukin 2 expression) /FL=gb:U12170.1	U12170		NP_110378
210878_s_at	5.03E-04	gb:BC001202.1 /DEF=Homo sapiens, clone MGC:3185, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3185) /DB_XREF=gi:12654720 /UG=Hs.24125 putative zinc finger protein /FL=gb:BC001202.1	BC001202		NP_057688
210879_s_at	0.002418	gb:AF334812.1 /DEF=Homo sapiens Rab11 interacting protein Rip11a mRNA, complete cds. /FEA=mRNA /PROD=Rab11 interacting protein Rip11a /DB_XREF=gi:13377896 /UG=Hs.24557 KIAA0857 protein /FL=gb:AF334812.1	AF334812		NP_056285
210891_s_at	3.26E-04	gb:AF035737.1 /DEF=Homo sapiens general transcription factor 2-I (GTF2I) mRNA, complete cds. /FEA=mRNA /GEN=GTF2I /PROD=general transcription factor 2-I; alternativesplice product /DB_XREF=gi:2827179 /UG=Hs.278589 general transcription factor II, i /FL=gb:AF035737.1	AF035737		NP_127496

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210907_s_at	1.26E-04	gb:BC002506.1 /DEF=Homo sapiens, programmed cell death 10, clone MGC:1212, mRNA, complete cds. /FEA=mRNA /PROD=programmed cell death 10 /DB_XREF=gi:12803370 /UG=Hs.28866 programmed cell death 10 /FL=gb:BC002506.1	BC002506		NP_665859
210915_x_at	7.58E-04	gb:M15564.1 /DEF=Human T-cell receptor rearranged beta-chain V-region (V-D-J) mRNA, complete cds. /FEA=mRNA /GEN=TCRB /DB_XREF=gi:339011 /UG=Hs.303157 T cell receptor beta locus /FL=gb:M15564.1	M15564		
210926_at	0.00228	gb:AY014272.1 /DEF=Homo sapiens FKSG30 (FKSG30) mRNA, complete cds. /FEA=mRNA /GEN=FKSG30 /PROD=FKSG30 /DB_XREF=gi:12408251 /UG=Hs.315492 Homo sapiens FKSG30 (FKSG30) mRNA, complete cds /FL=gb:AY014272.1	AY014272		
210927_x_at	7.58E-04	gb:BC004239.1 /DEF=Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRNA, complete cds. /FEA=mRNA /PROD=jumping translocation breakpoint /DB_XREF=gi:13278986 /UG=Hs.323093 Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRNA, complete cds /FL=gb:BC004239.1	BC004239		NP_006685
210942_s_at	0.010205	gb:AB022918.1 /DEF=Homo sapiens mRNA for alpha2,3-sialyltransferase ST3Gal VI, complete cds. /FEA=mRNA /GEN=ST3Gal VI /PROD=alpha2,3-sialyltransferase ST3Gal VI /DB_XREF=gi:4827246 /UG=Hs.34578 alpha2,3-sialyltransferase /FL=gb:AB022918.1	AB022918		NP_006091

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210943_s_at	1.26E-04	gb:U84744.1 /DEF=Human Chediak-Higashi syndrome protein short isoform (LYST) mRNA, complete cds. /FEA=mRNA /GEN=LYST /PROD=Chediak-Higashi syndrome protein short isoform /DB_XREF=gi:2654473 /UG=Hs.36508 Chediak-Higashi syndrome 1 /FL=gb:U84744.1	U84744		NP_000072
210944_s_at	0.005852	gb:BC003169.1 /DEF=Homo sapiens, Similar to calpain 3, (p94), clone MGC:4403, mRNA, complete cds. /FEA=mRNA /PROD=Similar to calpain 3, (p94) /DB_XREF=gi:13111992 /UG=Hs.40300 calpain 3, (p94) /FL=gb:BC003169.1	BC003169		NP_775113
210950_s_at	1.26E-04	gb:BC003573.1 /DEF=Homo sapiens, farnesyl-diphosphate farnesyltransferase 1, clone MGC:2200, mRNA, complete cds. /FEA=mRNA /PROD=farnesyl-diphosphate farnesyltransferase 1 /DB_XREF=gi:13097746 /UG=Hs.48876 farnesyl-diphosphate farnesyltransferase 1 /FL=gb:BC003573.1	BC003573		NP_004453
210951_x_at	3.26E-04	gb:AF125393.1 /DEF=Homo sapiens Rab27 isoform mRNA, complete cds. /FEA=mRNA /PROD=Rab27 isoform /DB_XREF=gi:5410356 /UG=Hs.50477 RAB27A, member RAS oncogene family /FL=gb:AF125393.1	AF125393		NP_004571

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210965_x_at	0.002418	gb:BC001274.1 /DEF=Homo sapiens, Similar to cell division cycle 2-like 5 (cholinesterase-related cell division controller), clone MGC:5091, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cell division cycle 2-like 5(cholinesterase-related cell division controller) /DB_XREF=gi:12654860 /UG=Hs.59498 cell division cycle 2-like 5 (cholinesterase-related cell division controller) /FL=gb:BC001274.1	BC001274		NP_112557
210966_x_at	0.00545	gb:BC001460.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ10378, clone MGC:2954, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ10378 /DB_XREF=gi:12655204 /UG=Hs.6214 KIAA0731 protein /FL=gb:BC001460.1	BC001460		NP_056130
210968_s_at	0.02145	gb:AF333336.1 /DEF=Homo sapiens testis specific reticulon 5 protein mRNA, complete cds. /FEA=mRNA /PROD=testis specific reticulon 5 protein /DB_XREF=gi:13377627 /UG=Hs.65450 reticulon 4 /FL=gb:AF333336.1	AF333336		NP_722550
210969_at	0.003529	gb:AF118089.1 /DEF=Homo sapiens PRO2042 mRNA, complete cds. /FEA=mRNA /PROD=PRO2042 /DB_XREF=gi:6650823 /UG=Hs.69171 protein kinase C-like 2 /FL=gb:AF118089.1	AF118089		NP_006247
210972_x_at	2.05E-04	gb:M15565.1 /DEF=Human T-cell receptor rearranged alpha-chain V-region (V-D-J) mRNA, complete cds. /FEA=mRNA /GEN=TCRA /DB_XREF=gi:338765 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:M15565.1	M15565		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210980_s_at	1.26E-04	gb:U47674.1 /DEF=Homo sapiens putative heart protein PHP mRNA, complete cds. /FEA=mRNA /PROD=PHP /DB_XREF=gi:3860239 /UG=Hs.75811 N-acylsphingosine amidohydrolase (acid ceramidase) /FL=gb:U47674.1	U47674		NP_808592
210981_s_at	0.001618	gb:AF040751.1 /DEF=Homo sapiens G protein-coupled receptor kinase 6, splice variant B (GRK6) mRNA, complete cds. /FEA=mRNA /GEN=GRK6 /PROD=G protein-coupled receptor kinase 6, splicevariant B /DB_XREF=gi:3005015 /UG=Hs.76297 G protein-coupled receptor kinase 6 /FL=gb:AF040751.1	AF040751		NP_002073
210986_s_at	0.00161	gb:Z24727.1 /DEF=H.sapiens tropomyosin isoform mRNA, complete CDS. /FEA=mRNA /PROD=tropomyosin isoform /DB_XREF=gi:854188 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:Z24727.1	Z24727		NP_000357
210987_x_at	1.26E-04	gb:M19267.1 /DEF=Human tropomyosin mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:339943 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:M19267.1	M19267		
210996_s_at	3.26E-04	gb:U43430.1 /DEF=Human epsilon isoform 14-3-3 protein mRNA, complete cds. /FEA=mRNA /PROD=14-3-3 protein /DB_XREF=gi:4096984 /UG=Hs.79474 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, epsilon polypeptide /FL=gb:U43399.1 gb:U43430.1	U43430		NP_006752

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211015_s_at	2.05E-04	gb:L12723.1 /DEF=Human heat shock protein 70 (hsp70) mRNA, complete cds. /FEA=mRNA /GEN=hsp70 /PROD=heat shock protein 70 /DB_XREF=gi:292159 /UG=Hs.90093 heat shock 70kD protein 4 /FL=gb:L12723.1	L12723		
211025_x_at	1.26E-04	gb:BC006229.1 /DEF=Homo sapiens, cytochrome c oxidase subunit Vb, clone MGC:10622, mRNA, complete cds. /FEA=mRNA /PROD=cytochrome c oxidase subunit Vb /DB_XREF=gi:13623258 /FL=gb:BC006229.1	BC006229		NP_001853
211026_s_at	0.004341	gb:BC006230.1 /DEF=Homo sapiens, lysophospholipase-like, clone MGC:10338, mRNA, complete cds. /FEA=mRNA /PROD=lysophospholipase-like /DB_XREF=gi:13623260 /FL=gb:BC006230.1	BC006230		NP_009214
211034_s_at	0.005339	gb:BC006270.1 /DEF=Homo sapiens, clone MGC:11291, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:11291) /DB_XREF=gi:13623331 /FL=gb:BC006270.1	BC006270		
211040_x_at	0.004341	gb:BC006325.1 /DEF=Homo sapiens, G-2 and S-phase expressed 1, clone MGC:12560, mRNA, complete cds. /FEA=mRNA /PROD=G-2 and S-phase expressed 1 /DB_XREF=gi:13623450 /FL=gb:BC006325.1	BC006325		NP_057510
211043_s_at	0.001116	gb:BC006332.1 /DEF=Homo sapiens, clathrin, light polypeptide (Lcb), clone MGC:12930, mRNA, complete cds. /FEA=mRNA /PROD=clathrin, light polypeptide (Lcb) /DB_XREF=gi:13623458 /FL=gb:BC006332.1	BC006332		NP_009028

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211047_x_at	0.013223	gb:BC006337.1 /DEF=Homo sapiens, clone MGC:12798, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12798) /DB_XREF=gi:13623468 /FL=gb:BC006337.1	BC006337		NP_067586
211058_x_at	5.03E-04	gb:BC006379.1 /DEF=Homo sapiens, tubulin alpha 1, clone MGC:12832, mRNA, complete cds. /FEA=mRNA /PROD=tubulin alpha 1 /DB_XREF=gi:13623540 /FL=gb:BC006379.1	BC006379		NP_006073
211060_x_at	0.002418	gb:BC006383.1 /DEF=Homo sapiens, clone MGC:12663, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12663) /DB_XREF=gi:13623546 /FL=gb:BC006383.1	BC006383		NP_003792
211067_s_at	0.016074	gb:BC006454.1 /DEF=Homo sapiens, growth arrest-specific 7, clone MGC:1348, mRNA, complete cds. /FEA=mRNA /PROD=growth arrest-specific 7 /DB_XREF=gi:13623656 /FL=gb:BC006454.1	BC006454		NP_005881
211068_x_at	0.010205	gb:BC006456.1 /DEF=Homo sapiens, clone MGC:1426, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:1426) /DB_XREF=gi:13623660 /FL=gb:BC006456.1	BC006456		
211072_x_at	5.03E-04	gb:BC006481.1 /DEF=Homo sapiens, tubulin alpha 1, clone MGC:4387, mRNA, complete cds. /FEA=mRNA /PROD=tubulin alpha 1 /DB_XREF=gi:13623706 /FL=gb:BC006481.1	BC006481		NP_006073
211073_x_at	0.041055	gb:BC006483.1 /DEF=Homo sapiens, ribosomal protein L3, clone MGC:4393, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L3 /DB_XREF=gi:13623708 /FL=gb:BC006483.1	BC006483		NP_000958

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211074_at	0.003171	gb:AF000381.1 /DEF=Homo sapiens non-functional folate binding protein mRNA, complete cds. /FEA=mRNA /PROD=non-functional folate binding protein /DB_XREF=gi:2565195 /FL=gb:AF000381.1	AF000381		
211075_s_at	1.26E-04	gb:Z25521.1 /DEF=H.sapiens integrin associated protein mRNA, complete CDS,. /FEA=mRNA /PROD=integrin associated protein /DB_XREF=gi:396704 /FL=gb:Z25521.1	Z25521		NP_001768
211085_s_at	5.03E-04	gb:Z25430.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405738 /FL=gb:Z25430.1	Z25430		NP_006273
211087_x_at	0.011725	gb:Z25432.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405742 /FL=gb:Z25432.1	Z25432		NP_620583
211097_s_at	0.009689	gb:BC003111.1 /DEF=Homo sapiens, Similar to pre-B-cell leukemia transcription factor 2, clone MGC:2174, mRNA, complete cds. /FEA=mRNA /PROD=Similar to pre-B-cell leukemia transcriptionfactor 2 /DB_XREF=gi:13111886 /UG=Hs.93728 pre-B-cell leukemia transcription factor 2 /FL=gb:BC003111.1	BC003111		
211098_x_at	6.68E-04	gb:AF277194.1 /DEF=Homo sapiens PNAS-136 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-136 /DB_XREF=gi:12751112 /UG=Hs.93832 putative membrane protein /FL=gb:AF277194.1	AF277194		NP_061899

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211100_x_at	0.026891	gb:U82278.1 /DEF=Human immunoglobulin-like transcript 1c mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 1c /DB_XREF=gi:1907322 /UG=Hs.94498 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 /FL=gb:U82278.1 gb:AF025531.1	U82278		NP_006857
211121_s_at	0.002989	gb:AF180527.1 /DEF=Homo sapiens p22Dokdel (DOKDEL) mRNA, complete cds. /FEA=mRNA /GEN=DOKDEL /PROD=p22Dokdel /DB_XREF=gi:6606314 /UG=Hs.103854 docking protein 1, 62kD (downstream of tyrosine kinase 1) /FL=gb:AF180527.1	AF180527		NP_001372
211137_s_at	0.0375	gb:AF189723.2 /DEF=Homo sapiens calcium transport ATPase ATP2C1 (ATP2C1A) mRNA, complete cds. /FEA=mRNA /GEN=ATP2C1A /PROD=calcium transport ATPase ATP2C1 /DB_XREF=gi:6826913 /UG=Hs.106778 ATPase, Ca++ transporting, type 2C, member 1 /FL=gb:AF189723.2	AF189723		NP_055197
211144_x_at	0.026891	gb:M30894.1 /DEF=Human T-cell receptor Ti rearranged gamma-chain mRNA V-J-C region, complete cds. /FEA=mRNA /GEN=CD3G /DB_XREF=gi:339406 /UG=Hs.112259 T cell receptor gamma locus /FL=gb:M30894.1	M30894		
211160_x_at	0.033491	gb:M95178.1 /DEF=Human non-muscle alpha-actinin mRNA, complete cds. /FEA=mRNA /GEN=ACTN1 /PROD=alpha-actinin /DB_XREF=gi:178051 /UG=Hs.119000 actinin, alpha 1 /FL=gb:M95178.1	M95178		NP_001093

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211202_s_at	0.013223	gb:AF087481.1 /DEF=Homo sapiens retinoblastoma binding protein 2 homolog 1 (RBBP2H1) mRNA, complete cds. /FEA=mRNA /GEN=RBBP2H1 /PROD=retinoblastoma binding protein 2 homolog 1 /DB_XREF=gi:4322487 /UG=Hs.143323 putative DNAchromatin binding motif /FL=gb:AF087481.1	AF087481		NP_006609
211220_s_at	0.038535	gb:BC005329.1 /DEF=Homo sapiens, Similar to heat shock transcription factor 2, clone MGC:12415, mRNA, complete cds. /FEA=mRNA /PROD=Similar to heat shock transcription factor 2 /DB_XREF=gi:13529106 /UG=Hs.158195 heat shock transcription factor 2 /FL=gb:BC005329.1	BC005329		NP_004497
211251_x_at	0.007052	gb:U78774.1 /DEF=Human NFY-C mRNA, complete cds. /FEA=mRNA /PROD=NFY-C /DB_XREF=gi:2327008 /UG=Hs.168157 nuclear transcription factor Y, gamma /FL=gb:U78774.1	U78774		NP_055038
211252_x_at	0.013575	gb:U36759.1 /DEF=Human pre-T cell receptor alpha-type chain precursor, mRNA, complete cds. /FEA=mRNA /PROD=pre-T cell receptor alpha-type chain precursor /DB_XREF=gi:1127580 /UG=Hs.169002 Human pre TCR alpha mRNA, partial cds /FL=gb:U36759.1	U36759		NP_612153
211255_x_at	0.004541	gb:AF064605.1 /DEF=Homo sapiens KE05 protein mRNA, complete cds. /FEA=mRNA /PROD=KE05 protein /DB_XREF=gi:3152663 /UG=Hs.169681 death effector domain-containing /FL=gb:AF064605.1	AF064605		NP_127491

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211275_s_at	1.26E-04	gb:AF087942.1 /DEF=Homo sapiens glycogenin-1L mRNA, complete cds. /FEA=mRNA /PROD=glycogenin-1L /DB_XREF=gi:5814084 /UG=Hs.174071 glycogenin /FL=gb:AF087942.1	AF087942		NP_004121
211284_s_at	0.041055	gb:BC000324.1 /DEF=Homo sapiens, Similar to granulin, clone MGC:8480, mRNA, complete cds. /FEA=mRNA /PROD=Similar to granulin /DB_XREF=gi:12653114 /UG=Hs.180577 granulin /FL=gb:BC000324.1	BC000324		NP_002078
211285_s_at	0.041055	gb:U84404.1 /DEF=Human E6-associated protein E6-APubiquitin-protein ligase (UBE3A) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=UBE3A /PROD=E6-associated protein E6-APubiquitin-proteinligase /DB_XREF=gi:1872513 /UG=Hs.180686 ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome) /FL=gb:AF273050.1 gb:U84404.1	U84404		NP_570854
211286_x_at	0.009716	gb:L29349.1 /DEF=Homo sapiens granulocyte-macrophage colony-stimulating factor receptor alpha-subunit 3 (GM-CSF-RA3) mRNA, complete cds. /FEA=mRNA /GEN=GM-CSF-RA3 /PROD=granulocyte-macrophage colony-stimulating factorreceptor alpha-subunit 3 /DB_XREF=gi:460284 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:L29349.1	L29349		NP_758452

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211287_x_at	1.26E-04	gb:M64445.1 /DEF=Human GM-CSF receptor mRNA, complete cds. /FEA=mRNA /GEN=GM-CSF receptor /PROD=GM-CSF receptor /DB_XREF=gi:183361 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:M64445.1	M64445		NP_758452
211296_x_at	0.026891	gb:AB009010.1 /DEF=Homo sapiens mRNA for polyubiquitin UbC, complete cds. /FEA=mRNA /GEN=UbC1 /PROD=polyubiquitin UbC /DB_XREF=gi:2647407 /UG=Hs.183704 ubiquitin C /FL=gb:BC000449.1 gb:AB009010.1	AB009010		NP_066289
211297_s_at	1.28E-04	gb:L20320.1 /DEF=Human protein serinethreonine kinase stk1 mRNA, complete cds. /FEA=mRNA /PROD=protein serinethreonine kinase /DB_XREF=gi:348242 /UG=Hs.184298 cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase) /FL=gb:L20320.1	L20320		NP_001790
211307_s_at	0.005276	gb:U43677.1 /DEF=Human Fc alpha receptor, splice variant FcalphaR a.3 (CD89) mRNA, complete cds. /FEA=mRNA /GEN=CD89 /PROD=Fc alpha receptor /DB_XREF=gi:1439557 /UG=Hs.193122 Fc fragment of IgA, receptor for /FL=gb:U43677.1 gb:D87854.1 gb:D87856.1	U43677		NP_579814
211318_s_at	2.05E-04	gb:U85943.1 /DEF=Homo sapiens mRNA-associated protein mrnp41 mRNA, complete cds. /FEA=mRNA /PROD=mRNA-associated protein mrnp41 /DB_XREF=gi:2231591 /UG=Hs.196209 RAE1 (RNA export 1, S.pombe) homolog /FL=gb:U85943.1	U85943		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211325_x_at	0.007779	gb:U72518.1 /DEF=Human destrin-2 pseudogene mRNA, complete cds. /FEA=mRNA /PROD=destrin-2 /DB_XREF=gi:1673523 /UG=Hs.199299 Human destrin-2 pseudogene mRNA, complete cds /FL=gb:U72518.1	U72518		
211342_x_at	0.02145	gb:BC004354.1 /DEF=Homo sapiens, trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit), clone MGC:3120, mRNA, complete cds. /FEA=mRNA /PROD=trinucleotide repeat containing 11(THR-associated protein, 230 kDa subunit) /DB_XREF=gi:13279313 /UG=Hs.211607 trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) /FL=gb:BC004354.1	BC004354		
211355_x_at	0.013274	gb:U52914.1 /DEF=Human B219OB receptor isoform HuB219.3 precursor mRNA, complete cds. /FEA=mRNA /PROD=B219OB receptor isoform HuB219.3 precursor /DB_XREF=gi:1279904 /UG=Hs.226627 leptin receptor /FL=gb:U52914.1	U52914		NP_002294
211383_s_at	2.05E-04	gb:AL136827.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F2427 (from clone DKFZp434F2427); complete cds. /FEA=mRNA /GEN=DKFZp434F2427 /PROD=hypothetical protein /DB_XREF=gi:6807664 /UG=Hs.27207 KIAA0982 protein /FL=gb:AL136827.1	AL136827		NP_054742
211404_s_at	1.26E-04	gb:BC004371.1 /DEF=Homo sapiens, clone MGC:10449, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10449) /DB_XREF=gi:13325115 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC004371.1	BC004371		NP_001633

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211416_x_at	0.033381	gb:L20492.1 /DEF=Human gamma-glutamyl transpeptidase mRNA, complete cds. /FEA=mRNA /PROD=gamma-glutamyl transpeptidase /DB_XREF=gi:306752 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:L20492.1	L20492		NP_842564
211417_x_at	1.26E-04	gb:L20493.1 /DEF=Human gamma-glutamyl transpeptidase mRNA, complete cds. /FEA=mRNA /PROD=gamma-glutamyl transpeptidase /DB_XREF=gi:306754 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:L20493.1	L20493		
211426_x_at	0.011725	gb:U40038.1 /DEF=Human GTP-binding protein alpha q subunit (GNAQ) mRNA, complete cds. /FEA=mRNA /GEN=GNAQ /PROD=GTP-binding protein alpha q subunit /DB_XREF=gi:1181670 /UG=Hs.296261 guanine nucleotide binding protein (G protein), q polypeptide /FL=gb:U40038.1	U40038		NP_002063
211429_s_at	0.041055	gb:AF119873.1 /DEF=Homo sapiens PRO2275 mRNA, complete cds. /FEA=mRNA /PROD=PRO2275 /DB_XREF=gi:7770182 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 /FL=gb:AF119873.1	AF119873		NP_000286
211433_x_at	0.00161	gb:AL583909.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761J197 (from clone DKFZp761J197); complete cds. /FEA=mRNA /GEN=DKFZp761J197 /PROD=hypothetical protein /DB_XREF=gi:13093772 /UG=Hs.301696 hypothetical protein FLJ11560 /FL=gb:AL583909.1	AL583909		NP_079458

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211454_x_at	5.03E-04	gb:AF336878.1 /DEF=Homo sapiens FKSG51 (FKSG51) mRNA, complete cds. /FEA=mRNA /GEN=FKSG51 /PROD=FKSG51 /DB_XREF=gi:13384184 /UG=Hs.326752 Homo sapiens FKSG51 (FKSG51) mRNA, complete cds /FL=gb:AF336878.1	AF336878		
211471_s_at	0.028067	gb:AF133588.1 /DEF=Homo sapiens RAB36 (RAB36) mRNA, complete cds. /FEA=mRNA /GEN=RAB36 /PROD=RAB36 /DB_XREF=gi:6049163 /UG=Hs.38772 RAB36, member RAS oncogene family /FL=gb:AF133588.1	AF133588		NP_004905
211474_s_at	0.005852	gb:BC004948.1 /DEF=Homo sapiens, clone MGC:10846, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10846) /DB_XREF=gi:13436313 /UG=Hs.41072 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6 /FL=gb:BC004948.1	BC004948		
211475_s_at	1.26E-04	gb:AF116273.1 /DEF=Homo sapiens Bag1 protein variant mRNA, complete cds. /FEA=mRNA /PROD=Bag1 protein variant /DB_XREF=gi:4583370 /UG=Hs.41714 BCL2-associated athanogene /FL=gb:AF116273.1	AF116273		NP_004314
211505_s_at	1.26E-04	gb:AL136601.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J1516 (from clone DKFZp564J1516); complete cds. /FEA=mRNA /GEN=DKFZp564J1516 /PROD=hypothetical protein /DB_XREF=gi:13276702 /UG=Hs.6113 staufer (Drosophila, RNA-binding protein) /FL=gb:AL136601.1	AL136601		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211509_s_at	1.26E-04	gb:AB015639.1 /DEF=Homo sapiens ASY mRNA, complete cds. /FEA=mRNA /GEN=ASY /DB_XREF=gi:5821139 /UG=Hs.65450 reticulon 4 /FL=gb:AB015639.1	AB015639		NP_722550
211529_x_at	0.005852	gb:M90684.1 /DEF=Human lymphocyte antigen (HLA-G2.1) mRNA, complete cds. /FEA=mRNA /GEN=HLA-G /PROD=b2 microglobulin /DB_XREF=gi:188467 /UG=Hs.73885 HLA-G histocompatibility antigen, class I, G /FL=gb:M90684.1	M90684		NP_002118
211542_x_at	0.041055	gb:BC004334.1 /DEF=Homo sapiens, ribosomal protein S10, clone MGC:10943, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein S10 /DB_XREF=gi:13279259 /UG=Hs.76230 ribosomal protein S10 /FL=gb:BC004334.1	BC004334		NP_001005
211546_x_at	1.36E-04	gb:L36674.1 /DEF=Human (clone 2-5) synuclein (NACP) mRNA, complete cds. /FEA=mRNA /GEN=NACP /PROD=synuclein /DB_XREF=gi:556211 /UG=Hs.76930 synuclein, alpha (non A4 component of amyloid precursor) /FL=gb:L36674.1	L36674		NP_009292
211548_s_at	0.045541	gb:J05594.1 /DEF=Homo sapiens NAD+-dependent 15-hydroxyprostaglandin dehydrogenase (PDGH) mRNA, complete cds. /FEA=mRNA /GEN=PGDH /PROD=NAD+-dependent 15-hydroxyprostaglandin dehydrogenase /DB_XREF=gi:1203981 /UG=Hs.77348 hydroxyprostaglandin dehydrogenase 15-(NAD) /FL=gb:J05594.1	J05594		NP_000851

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211569_s_at	0.011725	gb:AF001903.1 /DEF=Human 3-hydroxyacyl-CoA dehydrogenase, isoform 2 mRNA, complete cds. /FEA=mRNA /PROD=3-hydroxyacyl-CoA dehydrogenase, isoform 2 /DB_XREF=gi:2078328 /UG=Hs.8110 L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain /FL=gb:AF001903.1	AF001903		NP_005318
211571_s_at	0.026891	gb:D32039.1 /DEF=Human pgH3 mRNA for proteoglycan PG-M(V3), complete cds. /FEA=mRNA /GEN=pgH3 /PROD=proteoglycan PG-M(V3) /DB_XREF=gi:1008912 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican) /FL=gb:D32039.1	D32039		NP_004376
211574_s_at	0.027418	gb:D84105.1 /DEF=Human CD46 mRNA, complete cds. /FEA=mRNA /PROD=CD46 /DB_XREF=gi:1256700 /UG=Hs.83532 membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen) /FL=gb:D84105.1	D84105		NP_758871
211581_x_at	0.007779	gb:AF000426.1 /DEF=Homo sapiens LST1 mRNA, cLST1E splice variant, complete cds. /FEA=mRNA /GEN=LST1 /DB_XREF=gi:2145067 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:AF000426.1	AF000426		NP_009092
211582_x_at	0.013223	gb:AF000424.1 /DEF=Homo sapiens LST1 mRNA, cLST1C splice variant, complete cds. /FEA=mRNA /GEN=LST1 /DB_XREF=gi:2145063 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:AF000424.1	AF000424		NP_009092
211609_x_at	0.00161	gb:U51007.1 /DEF=Human 26S protease subunit S5a mRNA, complete cds. /FEA=mRNA /PROD=26S protease subunit S5a /DB_XREF=gi:1256400 /FL=gb:U51007.1	U51007		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211623_s_at	0.005852	gb:M30448.1 /DEF=Human casein kinase II beta subunit mRNA, complete cds. /FEA=mRNA /GEN=CSNK2B /DB_XREF=gi:181154 /FL=gb:M30448.1	M30448		
211654_x_at	0.041055	gb:M17565.1 /DEF=Human MHC class II DQ-beta associated with DRw6, DQw1 protein, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB_XREF=gi:188188 /FL=gb:M17565.1	M17565		NP_002114
211657_at	2.05E-04	gb:M18728.1 /DEF=Human nonspecific crossreacting antigen mRNA, complete cds. /FEA=mRNA /GEN=NCA; NCA; NCA /PROD=non-specific cross reacting antigen /DB_XREF=gi:189084 /FL=gb:M18728.1	M18728		NP_002474
211666_x_at	0.005852	gb:L22453.1 /DEF=Homo sapiens HIV-1 TAR RNA binding protein (TARBP-b) mRNA, complete cds. /FEA=mRNA /GEN=TARBP-b /DB_XREF=gi:347963 /FL=gb:L22453.1	L22453		NP_000958
211671_s_at	0.005852	gb:U01351.1 /DEF=Human glucocorticoid receptor alpha-2 mRNA, complete cds. /FEA=mRNA /PROD=glucocorticoid receptor alpha-2 /DB_XREF=gi:458656 /FL=gb:U01351.1	U01351		NP_000167
211676_s_at	1.26E-04	gb:AF056979.1 /DEF=Homo sapiens clone YAN1 interferon-gamma receptor mRNA, complete cds. /FEA=mRNA /PROD=interferon-gamma receptor /DB_XREF=gi:13562048 /FL=gb:AF056979.1	AF056979		NP_000407
211684_s_at	3.26E-04	gb:AF250307.1 /DEF=Homo sapiens cytoplasmic dynein intermediate chain 2C mRNA, complete cds. /FEA=mRNA /PROD=cytoplasmic dynein intermediate chain 2C /DB_XREF=gi:13649464 /FL=gb:AF250307.1	AF250307		NP_001369

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211685_s_at	0.007052	gb:AF251061.1 /DEF=Homo sapiens neurocalcin mRNA, complete cds. /FEA=mRNA /PROD=neurocalcin /DB_XREF=gi:13625183 /FL=gb:AF251061.1	AF251061		NP_114430
211693_at	0.009689	gb:AF343666.1 /DEF=Homo sapiens translocation associated fusion protein IRTA1IGA1 (IRTA1IGHA1) mRNA, complete cds. /FEA=mRNA /GEN=IRTA1IGHA1 /PROD=translocation associated fusion protein IRTA1IGA1 /DB_XREF=gi:13591717 /FL=gb:AF343666.1	AF343666		
211696_x_at	0.00228	gb:AF349114.1 /DEF=Homo sapiens beta globin chain variant (HBB) mRNA, complete cds. /FEA=mRNA /GEN=HBB /PROD=beta globin chain variant /DB_XREF=gi:13549111 /FL=gb:AF349114.1	AF349114		NP_000509
211699_x_at	1.28E-04	gb:AF349571.1 /DEF=Homo sapiens hemoglobin alpha-1 globin chain (HBA1) mRNA, complete cds. /FEA=mRNA /GEN=HBA1 /PROD=hemoglobin alpha-1 globin chain /DB_XREF=gi:13650073 /FL=gb:AF349571.1	AF349571		NP_000549
211702_s_at	1.26E-04	gb:AF350251.1 /DEF=Homo sapiens ubiquitin specific protease mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin specific protease /DB_XREF=gi:13560796 /FL=gb:AF350251.1	AF350251		NP_115971
211703_s_at	5.03E-04	gb:AF353990.1 /DEF=Homo sapiens beta-amyloid binding protein precursor (BBP) mRNA, complete cds. /FEA=mRNA /GEN=BBP /PROD=beta-amyloid binding protein precursor /DB_XREF=gi:13625458 /FL=gb:AF353990.1	AF353990		NP_114416

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211710_x_at	0.010205	gb:BC005817.1 /DEF=Homo sapiens, ribosomal protein L4, clone MGC:11073, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L4 /DB_XREF=gi:13543303 /FL=gb:BC005817.1	BC005817		NP_000959
211711_s_at	0.013223	gb:BC005821.1 /DEF=Homo sapiens, phosphatase and tensin homolog (mutated in multiple advanced cancers 1), clone MGC:11227, mRNA, complete cds. /FEA=mRNA /PROD=phosphatase and tensin homolog (mutated in multiple advanced cancers 1) /DB_XREF=gi:13543309 /FL=gb:BC005821.1	BC005821		NP_000305
211725_s_at	1.26E-04	gb:BC005884.1 /DEF=Homo sapiens, clone MGC:4736, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:4736) /DB_XREF=gi:13543452 /FL=gb:BC005884.1	BC005884		
211727_s_at	0.02145	gb:BC005895.1 /DEF=Homo sapiens, COX11 (yeast) homolog, cytochrome c oxidase assembly protein, clone MGC:14469, mRNA, complete cds. /FEA=mRNA /PROD=COX11 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:13543474 /FL=gb:BC005895.1	BC005895		NP_004366
211729_x_at	0.003171	gb:BC005902.1 /DEF=Homo sapiens, Similar to biliverdin reductase A, clone MGC:14491, mRNA, complete cds. /FEA=mRNA /PROD=Similar to biliverdin reductase A /DB_XREF=gi:13543488 /FL=gb:BC005902.1	BC005902		NP_000703

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211730_s_at	0.001116	gb:BC005903.1 /DEF=Homo sapiens, polymerase (RNA) II (DNA directed) polypeptide L (7.6kD), clone MGC:14494, mRNA, complete cds. /FEA=mRNA /PROD=polymerase (RNA) II (DNA directed) polypeptide L(7.6kD) /DB_XREF=gi:13543491 /FL=gb:BC005903.1	BC005903		NP_066951
211733_x_at	0.010205	gb:BC005911.1 /DEF=Homo sapiens, sterol carrier protein 2, clone MGC:14505, mRNA, complete cds. /FEA=mRNA /PROD=sterol carrier protein 2 /DB_XREF=gi:13543502 /FL=gb:BC005911.1	BC005911		NP_002970
211741_x_at	0.041795	gb:BC005924.1 /DEF=Homo sapiens, pregnancy specific beta-1-glycoprotein 3, clone MGC:14526, mRNA, complete cds. /FEA=mRNA /PROD=pregnancy specific beta-1-glycoprotein 3 /DB_XREF=gi:13543532 /FL=gb:BC005924.1	BC005924		NP_066296
211742_s_at	0.001116	gb:BC005926.1 /DEF=Homo sapiens, ecotropic viral integration site 2B, clone MGC:14529, mRNA, complete cds. /FEA=mRNA /PROD=ecotropic viral integration site 2B /DB_XREF=gi:13543535 /FL=gb:BC005926.1	BC005926		NP_006486
211744_s_at	1.26E-04	gb:BC005930.1 /DEF=Homo sapiens, Similar to CD58 antigen, (lymphocyte function-associated antigen 3), clone MGC:14538, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CD58 antigen, (lymphocytefunction-associated antigen 3) /DB_XREF=gi:13543544 /FL=gb:BC005930.1	BC005930		NP_001770

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211745_x_at	7.58E-04	gb:BC005931.1 /DEF=Homo sapiens, hemoglobin, alpha 2, clone MGC:14541, mRNA, complete cds. /FEA=mRNA /PROD=hemoglobin, alpha 2 /DB_XREF=gi:13543547 /FL=gb:BC005931.1	BC005931		NP_000508
211746_x_at	0.004341	gb:BC005932.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, alpha type, 1, clone MGC:14542, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) subunit, alphas type, 1 /DB_XREF=gi:13543550 /FL=gb:BC005932.1	BC005932		NP_683877
211749_s_at	0.026891	gb:BC005941.1 /DEF=Homo sapiens, Similar to vesicle-associated membrane protein 3, clone MGC:14563, mRNA, complete cds. /FEA=mRNA /PROD=Similar to vesicle-associated membrane protein3 /DB_XREF=gi:13543573 /FL=gb:BC005941.1	BC005941		NP_004772
211755_s_at	0.001116	gb:BC005960.1 /DEF=Homo sapiens, Similar to ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit b, isoform 1, clone MGC:14609, mRNA, complete cds. /FEA=mRNA /PROD=Similar to ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit b, isoform 1 /DB_XREF=gi:13543617 /FL=gb:BC005960.1	BC005960		NP_001679
211758_x_at	0.007779	gb:BC005968.1 /DEF=Homo sapiens, ATP binding protein associated with cell differentiation, clone MGC:14620, mRNA, complete cds. /FEA=mRNA /PROD=ATP binding protein associated with celldifferentiation /DB_XREF=gi:13543638 /FL=gb:BC005968.1	BC005968		NP_005774

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211760_s_at	1.26E-04	gb:BC005974.1 /DEF=Homo sapiens, vesicle-associated membrane protein 4, clone MGC:14658, mRNA, complete cds. /FEA=mRNA /PROD=vesicle-associated membrane protein 4 /DB_XREF=gi:13543647 /FL=gb:BC005974.1	BC005974		NP_003753
211763_s_at	7.58E-04	gb:BC005979.1 /DEF=Homo sapiens, ubiquitin-conjugating enzyme E2B (RAD6 homolog), clone MGC:14670, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin-conjugating enzyme E2B (RAD6 homolog) /DB_XREF=gi:13543659 /FL=gb:BC005979.1	BC005979		NP_003328
211764_s_at	0.003171	gb:BC005980.1 /DEF=Homo sapiens, ubiquitin-conjugating enzyme E2D 1 (homologous to yeast UBC45), clone MGC:14673, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin-conjugating enzyme E2D 1 (homologous to yeast UBC45) /DB_XREF=gi:13543662 /FL=gb:BC005980.1	BC005980		NP_003329
211776_s_at	0.00161	gb:BC006141.1 /DEF=Homo sapiens, Similar to erythrocyte protein band 4.1-like 3, clone MGC:13087, mRNA, complete cds. /FEA=mRNA /PROD=Similar to erythrocyte protein band 4.1-like 3 /DB_XREF=gi:13544008 /FL=gb:BC006141.1	BC006141		NP_036439
211779_x_at	1.26E-04	gb:BC006155.1 /DEF=Homo sapiens, clone MGC:13188, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13188) /DB_XREF=gi:13544040 /FL=gb:BC006155.1	BC006155		NP_036437

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211782_at	0.031213	gb:BC006170.1 /DEF=Homo sapiens, Similar to iduronate 2-sulfatase (Hunter syndrome), clone MGC:13238, mRNA, complete cds. /FEA=mRNA /PROD=Similar to iduronate 2-sulfatase (Huntersyndrome) /DB_XREF=gi:13544076 /FL=gb:BC006170.1	BC006170		NP_006114
211783_s_at	0.001116	gb:BC006177.1 /DEF=Homo sapiens, Similar to metastasis associated 1, clone MGC:13258, mRNA, complete cds. /FEA=mRNA /PROD=Similar to metastasis associated 1 /DB_XREF=gi:13544097 /FL=gb:BC006177.1	BC006177		NP_004680
211796_s_at	0.00161	gb:AF043179.1 /DEF=Homo sapiens T cell receptor beta chain (TCRBV13S1-TCRBJ2S1) mRNA, complete cds. /FEA=CDS /GEN=TCRBV13S1-TCRBJ2S1 /PROD=T cell receptor beta chain /DB_XREF=gi:3002924 /UG=Hs.303157 T cell receptor beta locus /FL=gb:AF043179.1	AF043179		
211801_x_at	0.044231	gb:AF329637.1 /DEF=Homo sapiens mitofusin 1 mRNA, nuclear gene for mitochondrial protein, complete cds. /FEA=CDS /PROD=mitofusin 1 precursor /DB_XREF=gi:12744895 /UG=Hs.197877 hypothetical protein FLJ20693 /FL=gb:AF329637.1	AF329637		NP_284941
211814_s_at	0.012993	gb:AF112857.1 /DEF=Homo sapiens cyclin E2 splice variant 1 mRNA, complete cds. /FEA=CDS /PROD=cyclin E2 splice variant 1 /DB_XREF=gi:4092787 /UG=Hs.30464 cyclin E2 /FL=gb:AF112857.1	AF112857		NP_477097

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211816_x_at	0.004075	gb:D87858.1 /DEF=Homo sapiens mRNA for CD89_U04, complete cds. /FEA=CDS /PROD=CD89_U04 /DB_XREF=gi:1561545 /UG=Hs.193122 Fc fragment of IgA, receptor for /FL=gb:D87858.1 gb:D87860.1	D87858		NP_579814
211822_s_at	0.008036	gb:AF229061.1 /DEF=Homo sapiens NAC-gamma splice variant (NAC) mRNA, complete cds, alternatively spliced. /FEA=CDS /GEN=NAC /PROD=NAC-gamma splice variant /DB_XREF=gi:12656108 /UG=Hs.104305 death effector filament-forming Ced-4-like apoptosis protein /FL=gb:AF229061.1	AF229061		NP_127500
211825_s_at	0.04734	gb:AF327066.1 /DEF=Homo sapiens Ewings sarcoma EWS-Fli1 (type 1) oncogene mRNA, complete cds. /FEA=CDS /PROD=Ewings sarcoma EWS-Fli1 (type 1) oncogene /DB_XREF=gi:12963354 /UG=Hs.129953 Ewing sarcoma breakpoint region 1 /FL=gb:AF327066.1	AF327066		
211858_x_at	1.26E-04	gb:AF088184.1 /DEF=Homo sapiens guanine nucleotide-binding protein Gs alpha subunit isoform L2 mRNA, complete cds. /FEA=CDS /PROD=guanine nucleotide-binding protein Gs alphasubunit isoform L2 /DB_XREF=gi:3552031 /UG=Hs.273385 guanine nucleotide binding protein (G protein), alpha stimulating activity polypeptide 1 /FL=gb:AF088184.1	AF088184		NP_536351

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211883_x_at	0.041055	gb:M76742.1 /DEF=Homo sapiens alternatively spliced biliary glycoprotein (BGP _a) mRNA, complete cds. /FEA=CDS /GEN=BGP _a /PROD=biliary glycoprotein /DB_XREF=gi:179480 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:M76742.1	M76742		NP_001703
211889_x_at	0.026891	gb:D12502.1 /DEF=Human mRNA for biliary glycoprotein, complete cds. /FEA=CDS /GEN=BGP /PROD=biliary glycoprotein /DB_XREF=gi:219494 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:D12502.1	D12502		NP_001703
211902_x_at	0.004341	gb:L34703.1 /DEF=Homo sapiens T-cell receptor alpha chain (TCRA) mRNA (HLA-A1, 24; B7, 8; DR 1, 3), complete cds. /FEA=CDS /GEN=TCRA /PROD=T-cell receptor alpha /DB_XREF=gi:1100165 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:L34703.1	L34703		
211909_x_at	0.046651	gb:L32662.1 /DEF=Human prostaglandin E2 receptor EP3 subtype isoform IV mRNA, complete cds. /FEA=CDS /PROD=prostaglandin E2 receptor /DB_XREF=gi:484163 /FL=gb:L32662.1	L32662		
211913_s_at	0.008391	gb:L08961.1 /DEF=Homo sapiens transmembrane tyrosine kinase mRNA, complete cds. /FEA=CDS /PROD=tyrosine kinase /DB_XREF=gi:897614 /FL=gb:L08961.1	L08961		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211921_x_at	0.033381	gb:AF348514.1 /DEF=Homo sapiens fetal thymus prothymosin alpha mRNA, complete cds. /FEA=CDS /PROD=prothymosin alpha /DB_XREF=gi:13560658 /FL=gb:AF348514.1	AF348514		NP_002814
211926_s_at	0.041055	myosin, heavy polypeptide 9, non-muscle	AI827941	Hs.146550	NP_002464
211927_x_at	0.02145	ESTs, Highly similar to S22655 translation elongation factor eEF-1 gamma chain - human [H.sapiens]	BE963164	Hs.381229	
211928_at	0.004341	Consensus includes gb:AB002323.1 /DEF=Human mRNA for KIAA0325 gene, partial cds. /FEA=mRNA /GEN=KIAA0325 /DB_XREF=gi:2224590 /UG=Hs.7720 dynein, cytoplasmic, heavy polypeptide 1	AB002323		NP_001367
211930_at	0.004341	Homo sapiens BX1 mRNA, partial cds	BE867771	Hs.249247	
211936_at	7.58E-04	Consensus includes gb:AF216292.1 /DEF=Homo sapiens endoplasmic reticulum luminal Ca2+ binding protein grp78 mRNA, complete cds. /FEA=CDS /PROD=endoplasmic reticulum luminal Ca2+ bindingprotein grp78 /DB_XREF=gi:7229461 /UG=Hs.75410 heat shock 70kD protein 5 (glucose-regulated protein, 78kD) /FL=gb:AF216292.1	AF216292		NP_005338
211937_at	0.003171	Consensus includes gb:NM_001417.1 /DEF=Homo sapiens eukaryotic translation initiation factor 4B (EIF4B), mRNA. /FEA=CDS /GEN=EIF4B /PROD=eukaryotic translation initiation factor 4B /DB_XREF=gi:4503532 /UG=Hs.93379 eukaryotic translation initiation factor 4B /FL=gb:NM_001417.1	NM_001417		NP_001408
211941_s_at	0.033381	prostatic binding protein	BF686267	Hs.80423	NP_002558

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211944_at	0.003539	Consensus includes gb:BE729523 /FEA=EST /DB_XREF=gi:10143515 /DB_XREF=est:601564896F1 /CLONE=IMAGE:3926279 /UG=Hs.69559 KIAA1096 protein	AL096857		NP_055987
211946_s_at	0.026891	Consensus includes gb:AL096857.1 /DEF=Novel human mRNA from chromosome 1, which has similarities to BAT2 genes. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:5541862 /UG=Hs.69559 KIAA1096 protein	AL096857		NP_055987
211948_x_at	6.68E-04	Consensus includes gb:BG261071 /FEA=EST /DB_XREF=gi:12770887 /DB_XREF=est:602372693F1 /CLONE=IMAGE:4480631 /UG=Hs.69559 KIAA1096 protein	AL096857		NP_055987
211950_at	0.041055	Consensus includes gb:AB007931.1 /DEF=Homo sapiens mRNA for KIAA0462 protein, partial cds. /FEA=mRNA /GEN=KIAA0462 /PROD=KIAA0462 protein /DB_XREF=gi:3413885 /UG=Hs.297641 KIAA0462 protein	AB007931		NP_065816
211960_s_at	1.26E-04	Consensus includes gb:BG261416 /FEA=EST /DB_XREF=gi:12771232 /DB_XREF=est:602373192F1 /CLONE=IMAGE:4484422 /UG=Hs.237955 hypothetical protein PRO2706	AK000826		NP_004628
211961_s_at	1.26E-04	Consensus includes gb:AK000826.1 /DEF=Homo sapiens cDNA FLJ20819 fis, clone ADSE00511. /FEA=mRNA /DB_XREF=gi:7021139 /UG=Hs.237955 hypothetical protein PRO2706	AK000826		NP_004628
211963_s_at	5.03E-04	actin related protein 2/3 complex, subunit 5, 16kDa	AL516350	Hs.82425	NP_005708

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211967_at	0.026891	pro-oncosis receptor inducing membrane injury gene	BG538627	Hs.393431	NP_443164
211974_x_at	0.026891	recombining binding protein suppressor of hairless (Drosophila)	AL513759	Hs.356806	
211976_at	5.03E-04	Consensus includes gb:AK026168.1 /DEF=Homo sapiens cDNA: FLJ22515 fis, clone HRC12122, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence. /FEA=mRNA /DB_XREF=gi:10438931 /UG=Hs.188882 Homo sapiens cDNA: FLJ21862 fis, clone HEP02321, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence	AK026168		
211984_at	0.00161	ESTs, Moderately similar to apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2; apolipoprotein B mRNA editing enzyme, catalytic polypeptide 2 [Homo sapiens] [H.sapiens]	AI653730	Hs.374441	
211985_s_at	5.03E-04	ESTs, Moderately similar to apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2; apolipoprotein B mRNA editing enzyme, catalytic polypeptide 2 [Homo sapiens] [H.sapiens]	AI653730	Hs.374441	
211993_at	0.027792	protein kinase, lysine deficient 1	AI742553	Hs.184592	NP_061852
211994_at	1.26E-04	protein kinase, lysine deficient 1	AI742553	Hs.184592	NP_061852
211997_x_at	0.033381	Consensus includes gb:NM_005324.1 /DEF=Homo sapiens H3 histone, family 3B (H3.3B) (H3F3B), mRNA. /FEA=CDS /GEN=H3F3B /PROD=H3 histone, family 3B (H3.3B) /DB_XREF=gi:4885384 /UG=Hs.180877 H3 histone, family 3B (H3.3B) /FL=gb:NM_005324.1	NM_005324		NP_005315
212006_at	0.033381	Consensus includes gb:AU149908 /FEA=EST /DB_XREF=gi:11011429 /DB_XREF=est:AU149908 /CLONE=NT2RP2000896 /UG=Hs.77495 UBX domain-containing 1	D87684		NP_055422

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212007_at	0.011626	Consensus includes gb:A1927512 /FEA=EST /DB_XREF=gi:5663476 /DB_XREF=est:wo90g11.x1 /CLONE=IMAGE:2462660 /UG=Hs.77495 UBX domain-containing 1	D87684		NP_055422
212010_s_at	0.001116	Consensus includes gb:AK025647.1 /DEF=Homo sapiens cDNA: FLJ21994 fis, clone HEP06577, highly similar to AF103803 Homo sapiens clone H41 unknown mRNA. /FEA=mRNA /DB_XREF=gi:10438231 /UG=Hs.283690 hypothetical protein /FL=gb:NM_017548.1	AK025647		NP_060018
212016_s_at	0.020793	polypyrimidine tract binding protein 1	AA679988	Hs.172550	NP_787041
212017_at	0.026891	ESTs, Weakly similar to S57447 HPBR11-7 protein - human [H.sapiens]	BF677404	Hs.409180	
212020_s_at	0.002537	antigen identified by monoclonal antibody Ki-67	BF001806	Hs.80976	NP_002408
212024_x_at	2.17E-04	see GenBank Accession Number U01184 for cDNA; similar to Drosophila melanogaster flil in GenBank Accession Number U01182 and Caenorhabditis elegans flil homolog in GenBank Accession Number U01183; Homo sapiens FLII gene, complete cds.	NM_002018		NP_002009
212025_s_at	0.001054	see GenBank Accession Number U01184 for cDNA; similar to Drosophila melanogaster flil in GenBank Accession Number U01182 and Caenorhabditis elegans flil homolog in GenBank Accession Number U01183; Homo sapiens FLII gene, complete cds.	NM_002018		NP_002009
212027_at	1.26E-04	Homo sapiens cDNA: FLJ22454 fis, clone HRC09703	BE466128	Hs.409075	
212034_s_at	0.004341	likely ortholog of mouse exocyst component protein 70 kDa homolog (S. cerevisiae) Exo70: exocyst component protein 70 kDa homolog (S. cerevisiae)	BE646386	Hs.325530	NP_056034
212039_x_at	0.033381	ribosomal protein L3	BG339228	Hs.119598	NP_000958

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212040_at	0.033381	Consensus includes gb:BG249599 /FEA=EST /DB_XREF=gi:12759426 /DB_XREF=est:602319692F1 /CLONE=IMAGE:4414895 /UG=Hs.110771 Homo sapiens cDNA: FLJ21904 fis, clone HEP03585	AK025557		
212043_at	3.26E-04	Consensus includes gb:W72053 /FEA=EST /DB_XREF=gi:1382323 /DB_XREF=est:zd66e04.s1 /CLONE=IMAGE:345630 /UG=Hs.110771 Homo sapiens cDNA: FLJ21904 fis, clone HEP03585	AK025557		
212044_s_at	0.001116	ribosomal protein L27a	BE737027	Hs.76064	NP_000981
212046_x_at	0.041055	Consensus includes gb:X60188.1 /DEF=Human ERK1 mRNA for protein serinethreonine kinase. /FEA=mRNA /GEN=ERK1 /PROD=protein serinethreonine kinase /DB_XREF=gi:31220 /UG=Hs.861 mitogen-activated protein kinase 3	X60188		
212051_at	2.05E-04	Consensus includes gb:AA676803 /FEA=EST /DB_XREF=gi:2657325 /DB_XREF=est:zj65b04.s1 /CLONE=IMAGE:455119 /UG=Hs.13996 Homo sapiens cDNA: FLJ23260 fis, clone COL05804, highly similar to HSU90911 Human clone 23652 mRNA sequence	AK026913		NP_573571
212052_s_at	0.026891	Consensus includes gb:AB014576.1 /DEF=Homo sapiens mRNA for KIAA0676 protein, partial cds. /FEA=mRNA /GEN=KIAA0676 /PROD=KIAA0676 protein /DB_XREF=gi:3327165 /UG=Hs.155829 KIAA0676 protein	AB014576		NP_055858
212055_at	1.26E-04	DKFZP586M1523 protein	BF689173	Hs.22981	NP_056291

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212057_at	1.26E-04	Consensus includes gb:AA206161 /FEA=EST /DB_XREF=gi:1801720 /DB_XREF=est:zq55f01.s1 /CLONE=IMAGE:645529 /UG=Hs.75909 KIAA0182 protein	D80004		
212060_at	0.041055	Consensus includes gb:AU152088 /FEA=EST /DB_XREF=gi:11013609 /DB_XREF=est:AU152088 /CLONE=NT2RP3000162 /UG=Hs.7976 KIAA0332 protein	AB002330		
212063_at	5.03E-04	ESTs, Weakly similar to TRHY_HUMAN Trichohyalin [H.sapiens]	BE903880	Hs.408878	NP_000601
212065_s_at	0.003395	Consensus includes gb:AW502434 /FEA=EST /DB_XREF=gi:7116911 /DB_XREF=est:UI-HF-BR0p-ajs-g-03-0-UI.r1 /CLONE=IMAGE:3075652 /UG=Hs.180948 KIAA0729 protein	AB018272		
212066_s_at	0.013223	Consensus includes gb:AB018272.1 /DEF=Homo sapiens mRNA for KIAA0729 protein, partial cds. /FEA=mRNA /GEN=KIAA0729 /PROD=KIAA0729 protein /DB_XREF=gi:3882178 /UG=Hs.180948 KIAA0729 protein	AB018272		
212071_s_at	7.58E-04	spectrin, beta, non-erythrocytic 1	BE968833	Hs.107164	NP_842565
212072_s_at	0.001116	casein kinase 2, alpha 1 polypeptide pseudogene	AI161318	Hs.375587	
212076_at	0.011626	Consensus includes gb:AI701430 /FEA=EST /DB_XREF=gi:4989330 /DB_XREF=est:we29h08.x1 /CLONE=IMAGE:2342559 /UG=Hs.199160 myeloidlymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog) /FL=gb:L04284.1 gb:NM_005933.1	NM_005933		NP_005924

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212080_at	0.022795	Consensus includes gb:AV714029 /FEA=EST /DB_XREF=gi:10795546 /DB_XREF=est:AV714029 /CLONE=DCBCDA03 /UG=Hs.199160 myeloidlymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog) /FL=gb:L04284.1 gb:NM_005933.1	NM_005933		NP_005924
212082_s_at	0.00228	myosin, light polypeptide 6, alkali, smooth muscle and non-muscle	BE734356	Hs.77385	NP_524149
212086_x_at	0.016686	Consensus includes gb:AK026584.1 /DEF=Homo sapiens cDNA: FLJ22931 fis, clone KAT07501, highly similar to HSLAMAR Human mRNA for nuclear envelope protein lamin A precursor. /FEA=mRNA /DB_XREF=gi:10439468 /UG=Hs.77886 lamin AC	M13452		NP_733822
212098_at	0.00228	ESTs	AL134724	Hs.408961	
212099_at	0.041055	ras homolog gene family, member B	AI263909	Hs.204354	
212101_at	0.013575	Consensus includes gb:AU154321 /FEA=EST /DB_XREF=gi:11015842 /DB_XREF=est:AU154321 /CLONE=NT2RP4000774 /UG=Hs.301553 karyopherin alpha 6 (importin alpha 7) /FL=gb:AF060543.1 gb:NM_012316.1	NM_012316		NP_036448
212102_s_at	0.013223	Consensus includes gb:AK002111.1 /DEF=Homo sapiens cDNA FLJ11249 fis, clone PLACE1008790, highly similar to Homo sapiens importin alpha 7 subunit mRNA. /FEA=mRNA /DB_XREF=gi:7023792 /UG=Hs.301553 karyopherin alpha 6 (importin alpha 7) /FL=gb:AF060543.1 gb:NM_012316.1	NM_012316		NP_036448

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212112_s_at	0.003171	Consensus includes gb:AI816243 /FEA=EST /DB_XREF=gi:5431789 /DB_XREF=est:au46a11.x1 /CLONE=IMAGE:2517788 /UG=Hs.106823 H.sapiens gene from PAC 426I6, similar to syntaxin 7	AL035306		NP_803173
212120_at	1.26E-04	ras-like protein TC10	BF348067	Hs.250697	NP_036381
212124_at	5.03E-04	Consensus includes gb:AF070622.1 /DEF=Homo sapiens clone 24800 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283888 /UG=Hs.7252 KIAA1224 protein	AF070622		
212132_at	0.016934	Consensus includes gb:AL117499.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D1335 (from clone DKFZp434D1335); partial cds. /FEA=mRNA /GEN=DKFZp434D1335 /PROD=hypothetical protein /DB_XREF=gi:5912003 /UG=Hs.8258 DKFZP434D1335 protein	AL117499		NP_056393
212135_s_at	7.58E-04	ribosomal protein L35a	AW517686	Hs.287361	NP_001675
212136_at	0.005852	ribosomal protein L35a	AW517686	Hs.287361	NP_001675
212137_at	5.03E-04	hypothetical protein MGC19556	AV746402	Hs.356357	NP_291029
212150_at	0.033381	KIAA0143 protein	AA805651	Hs.84087	
212153_at	0.038535	Consensus includes gb:AB007930.1 /DEF=Homo sapiens mRNA for KIAA0461 peroteine, partial cds. /FEA=mRNA /GEN=KIAA0461 /PROD=KIAA0461 peroteine /DB_XREF=gi:3413883 /UG=Hs.107088 KIAA0461 protein	AB007930		NP_665739
212159_x_at	1.26E-04	Consensus includes gb:AI125280 /FEA=EST /DB_XREF=gi:3593794 /DB_XREF=est:qd92c05.x1 /CLONE=IMAGE:1736936 /UG=Hs.19121 adaptor-related protein complex 2, alpha 2 subunit	AB020706		NP_036437

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212162_at	0.026891	Consensus includes gb:AK022873.1 /DEF=Homo sapiens cDNA FLJ12811 fis, clone NT2RP2002475. /FEA=mRNA /DB_XREF=gi:10434518 /UG=Hs.9873 likely homolog of rat kinase D-interacting substance of 220 kDa; KIAA1250 protein	AB033076		
212163_at	0.004341	Consensus includes gb:AB033076.1 /DEF=Homo sapiens mRNA for KIAA1250 protein, partial cds. /FEA=mRNA /GEN=KIAA1250 /PROD=KIAA1250 protein /DB_XREF=gi:6330860 /UG=Hs.9873 likely homolog of rat kinase D-interacting substance of 220 kDa; KIAA1250 protein	AB033076		
212165_at	0.003171	Consensus includes gb:AF070537.1 /DEF=Homo sapiens clone 24606 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387894 /UG=Hs.17481 Homo sapiens clone 24606 mRNA sequence	AL133052		NP_612400
212166_at	2.76E-04	RAN binding protein 16	H38643	Hs.172685	NP_055839
212168_at	1.26E-04	Consensus includes gb:AL514547 /FEA=EST /DB_XREF=gi:12778041 /DB_XREF=est:AL514547 /CLONE=CL0BB004ZC07 (3 prime) /UG=Hs.180895 putative brain nuclearly-targeted protein	AB018308		NP_690051
212171_x_at	0.016426	vascular endothelial growth factor	H95344	Hs.73793	NP_003367
212175_s_at	0.007779	Consensus includes gb:AL513611 /FEA=EST /DB_XREF=gi:12777105 /DB_XREF=est:AL513611 /CLONE=XCL0BB001ZB07 (3 prime) /UG=Hs.171811 adenylate kinase 2	AK023758		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212177_at	0.003171	Consensus includes gb:AW081113 /FEA=EST /DB_XREF=gi:6036265 /DB_XREF=est:xc29c08.x1 /CLONE=IMAGE:2585678 /UG=Hs.18368 DKFZP564B0769 protein	AL080186		NP_116259
212180_at	1.26E-04	Consensus includes gb:AK000311.1 /DEF=Homo sapiens cDNA FLJ20304 fis, clone HEP06749. /FEA=mRNA /DB_XREF=gi:7020310 /UG=Hs.5613 Homo sapiens mRNA; cDNA DKFZp564E2222 (from clone DKFZp564E2222)	AK000311		NP_005198
212184_s_at	0.001116	Consensus includes gb:AL117407.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2050 (from clone DKFZp434D2050); partial cds. /FEA=mRNA /GEN=DKFZp434D2050 /PROD=hypothetical protein /DB_XREF=gi:591:1992 /UG=Hs.109727 TAK1-binding protein 2; KIAA0733 protein	AL117407		NP_663317
212191_x_at	0.026891	ribosomal protein L13	AW574664	Hs.180842	NP_150254
212192_at	0.004341	Consensus includes gb:AI718937 /FEA=EST /DB_XREF=gi:5036193 /DB_XREF=est:as50b04.x1 /CLONE=IMAGE:2320591 /UG=Hs.109438 Homo sapiens clone 24775 mRNA sequence	AF052169		NP_612453
212193_s_at	0.004644	likely ortholog of mouse la related protein	BE881529	Hs.6214	NP_056130
212198_s_at	5.03E-04	Consensus includes gb:AL515964 /FEA=EST /DB_XREF=gi:12779457 /DB_XREF=est:AL515964 /CLONE=CS0DA001YL22 (3 prime) /UG=Hs.79305 KIAA0255 gene product	AL049539		
212200_at	0.016934	Consensus includes gb:AK025933.1 /DEF=Homo sapiens cDNA: FLJ22280 fis, clone HRC03841. /FEA=mRNA /DB_XREF=gi:10438600 /UG=Hs.100729 KIAA0692 protein	AB014592		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212202_s_at	0.00228	Consensus includes gb:BG493972 /FEA=EST /DB_XREF=gi:13455486 /DB_XREF=est:602542252F1 /CLONE=IMAGE:4673316 /UG=Hs.16492 DKFZP564G2022 protein	AF132733		NP_056312
212204_at	2.05E-04	Consensus includes gb:AL049944.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G2022 (from clone DKFZp564G2022); partial cds. /FEA=mRNA /GEN=DKFZp564G2022 /PROD=hypothetical protein /DB_XREF=gi:4884189 /UG=Hs.16492 DKFZP564G2022 protein	AF132733		NP_056312
212208_at	7.58E-04	Consensus includes gb:AK023837.1 /DEF=Homo sapiens cDNA FLJ13775 fis, clone PLACE4000369, moderately similar to Homo sapiens thyroid hormone receptor-associated protein complex component TRAP240 mRNA. /FEA=mRNA /DB_XREF=gi:10435894 /UG=Hs.4084 KIAA1025 protein	AK023837		
212211_at	1.26E-04	gene trap ankyrin repeat	AI986295	Hs.12329	NP_115593
212219_at	0.003171	Consensus includes gb:D38521.1 /DEF=Human mRNA for KIAA0077 gene, partial cds. /FEA=mRNA /GEN=KIAA0077 /DB_XREF=gi:559329 /UG=Hs.112396 KIAA0077 protein	D38521		
212223_at	0.016934	Consensus includes gb:AI926544 /FEA=EST /DB_XREF=gi:5662508 /DB_XREF=est:wo46c12.x1 /CLONE=IMAGE:2458390 /UG=Hs.303154 popeye protein 3	AL117536		
212225_at	0.035763	putative translation initiation factor	AL516854	Hs.150580	NP_005792

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212228_s_at	0.012555	Consensus includes gb:AC004382 /DEF=Homo sapiens Chromosome 16 BAC clone CIT987SK-A-152E5 /FEA=mRNA_8 /DB_XREF=gi:3252819 /UG=Hs.4288 hypothetical protein DKFZp434K046	AC004382		
212229_s_at	6.68E-04	Consensus includes gb:AK001699.1 /DEF=Homo sapiens cDNA FLJ10837 fis, clone NT2RP4001260, highly similar to Homo sapiens mRNA for KIAA0875 protein. /FEA=mRNA /DB_XREF=gi:7023121 /UG=Hs.184227 F-box only protein 21 /FL=gb:AF174601.1	AK001699		NP_296373
212230_at	0.014233	ribosomal protein S20	AL576654	Hs.8102	NP_001014
212239_at	0.033381	Consensus includes gb:AI680192 /FEA=EST /DB_XREF=gi:4890374 /DB_XREF=est:tw66a10.x1 /CLONE=IMAGE:2264634 /UG=Hs.6241 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	M61906		NP_852556
212240_s_at	2.05E-04	Consensus includes gb:AI679268 /FEA=EST /DB_XREF=gi:4889450 /DB_XREF=est:tu62e04.x1 /CLONE=IMAGE:2255646 /UG=Hs.6241 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	M61906		NP_852556
212242_at	3.26E-04	tubulin, alpha 1 (testis specific)	AL565074	Hs.75318	NP_005991
212248_at	0.00228	Homo sapiens LYRIC mRNA, complete cds	AI972475	Hs.395896	
212249_at	0.008036	Consensus includes gb:AI934473 /FEA=EST /DB_XREF=gi:5673433 /DB_XREF=est:wp58d05.x1 /CLONE=IMAGE:2465961 /UG=Hs.6241 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	M61906		NP_852556
212250_at	1.26E-04	Homo sapiens LYRIC mRNA, complete cds	AI972475	Hs.395896	
212252_at	0.02145	calcium/calmodulin-dependent protein kinase kinase 2, beta	AA181179	Hs.108708	NP_757380

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212256_at	5.03E-04	putative UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T9	BE906572	Hs.394578	NP_060010
212259_s_at	0.001618	hematopoietic PBX-interacting protein	BF344265	Hs.8068	NP_065385
212261_at	0.001116	trinucleotide repeat containing 15	AL045800	Hs.323317	
212262_at	1.26E-04	Consensus includes gb:AA149639 /FEA=EST /DB_XREF=gi:1720440 /DB_XREF=est:zl39c06.s1 /CLONE=IMAGE:504298 /UG=Hs.15020 homolog of mouse quaking QKI (KH domain RNA binding protein) /FL=gb:AF142419.1 gb:AF142422.1	AF142419		
212264_s_at	3.26E-04	Consensus includes gb:BE645850 /FEA=EST /DB_XREF=gi:9970161 /DB_XREF=est:7e77c03.x1 /CLONE=IMAGE:3288484 /UG=Hs.154978 KIAA0261 protein	D87450		
212265_at	5.03E-04	Consensus includes gb:AL031781 /DEF=Human DNA sequence from clone 51J12 on chromosome 6q26-27. Contains the 3 part of the alternatively spliced gene for the human orthologs of mouse QKI-7 and QKI-7B (KH Domain RNA Binding proteins) and zebrafish ZKQ-1 (Quaking protein homolog). Con... /FEA=mRNA_1 /DB_XREF=gi:4038570 /UG=Hs.15020 homolog of mouse quaking QKI (KH domain RNA binding protein) /FL=gb:AF142419.1 gb:AF142422.1	AF142419		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212268_at	1.26E-04	Consensus includes gb:NM_030666.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 (SERPINB1), mRNA. /FEA=CDS /GEN=SERPINB1 /PROD=serine (or cysteine) proteinase inhibitor, cladeB (ovalbumin), member 1 /DB_XREF=gi:13489086 /UG=Hs.183583 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 /FL=gb:NM_030666.1	NM_030666		NP_109591
212271_at	1.26E-04	mitogen-activated protein kinase 1	AA195999	Hs.324473	NP_620407
212273_x_at	3.26E-04	GNAS complex locus	AI591100	Hs.374523	NP_536351
212277_at	0.003171	Consensus includes gb:AB014547.1 /DEF=Homo sapiens mRNA for KIAA0647 protein, partial cds. /FEA=mRNA /GEN=KIAA0647 /PROD=KIAA0647 protein /DB_XREF=gi:3327107 /UG=Hs.141727 myotubularin related protein 4 /FL=gb:NM_004687.1 gb:AF264717.1	AB014547		NP_004678
212293_at	0.010205	Consensus includes gb:BG111260 /FEA=EST /DB_XREF=gi:12604766 /DB_XREF=est:602281701F1 /CLONE=IMAGE:4369057 /UG=Hs.12259 KIAA0630 protein	AB014530		NP_689909
212297_at	1.26E-04	hypothetical protein FLJ20986	BF218804	Hs.324507	NP_078800
212301_at	1.26E-04	Consensus includes gb:D87440.1 /DEF=Human mRNA for KIAA0252 gene, partial cds. /FEA=mRNA /GEN=KIAA0252 /DB_XREF=gi:1665770 /UG=Hs.83419 KIAA0252 protein	D87440		NP_055953

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212306_at	0.041055	Consensus includes gb:AI741784 /FEA=EST /DB_XREF=gi:5110072 /DB_XREF=est:wg22h09.x1 /CLONE=IMAGE:2365889 /UG=Hs.108614 KIAA0627 protein; Drosophila multiple asters (Mast)-like homolog 2	AB014527		
212312_at	1.32E-04	Consensus includes gb:AL117381 /DEF=Human DNA sequence from clone RP5-857M17 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 3 part of the ID1 gene for inhibitor of DNA binding 1 (dominant negative helix-loop-helix protein), a gene for a novel protein simil... /FEA=mRNA_1 /DB_XREF=gi:11493211 /UG=Hs.305890 BCL2-like 1	AL117381		
212314_at	0.041055	Consensus includes gb:AB018289.1 /DEF=Homo sapiens mRNA for KIAA0746 protein, partial cds. /FEA=mRNA /GEN=KIAA0746 /PROD=KIAA0746 protein /DB_XREF=gi:3882212 /UG=Hs.49500 KIAA0746 protein	AB018289		
212315_s_at	0.023856	Consensus includes gb:AA502912 /FEA=EST /DB_XREF=gi:2237879 /DB_XREF=est:ne42d10.s1 /CLONE=IMAGE:900019 /UG=Hs.56966 KIAA0906 protein	AB020713		NP_079199
212317_at	0.005852	Consensus includes gb:AK022910.1 /DEF=Homo sapiens cDNA FLJ12848 fis, clone NT2RP2003391, highly similar to Homo sapiens mRNA for nuclear transport receptor. /FEA=mRNA /DB_XREF=gi:10434575 /UG=Hs.69235 transportin-SR /FL=gb:NM_012470.1	NM_012470		NP_036602

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212320_at	0.02145	Consensus includes gb:BC001002.1 /DEF=Homo sapiens, clone IMAGE:3447696, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3447696) /DB_XREF=gi:12654358 /UG=Hs.179661 tubulin, beta polypeptide	BC001002		NP_821133
212322_at	0.00161	Consensus includes gb:BE999972 /FEA=EST /DB_XREF=gi:10700248 /DB_XREF=est:7h15b02.x1 /CLONE=IMAGE:3316011 /UG=Hs.186613 sphingosine-1-phosphate lyase 1 /FL=gb:AF144638.1	AF144638		NP_003892
212323_s_at	3.26E-04	Consensus includes gb:AA524553 /FEA=EST /DB_XREF=gi:2265481 /DB_XREF=est:ng45h12.s1 /CLONE=IMAGE:937799 /UG=Hs.194737 KIAA0453 protein	AB007922		
212330_at	1.26E-04	transcription factor Dp-1	R60866	Hs.79353	NP_009042
212332_at	0.020793	Consensus includes gb:BF110947 /FEA=EST /DB_XREF=gi:10940637 /DB_XREF=est:7n30f02.x1 /CLONE=IMAGE:3566330 /UG=Hs.79362 retinoblastoma-like 2 (p130) /FL=gb:NM_005611.1	NM_005611		NP_005602
212334_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW167793	Hs.409152	NP_002067
212335_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW167793	Hs.409152	NP_002067
212340_at	2.05E-04	general transcription factor IIA, 2, 12kDa	AA195936	Hs.76362	
212341_at	0.005852	general transcription factor IIA, 2, 12kDa	AA195936	Hs.76362	
212342_at	0.011725	general transcription factor IIA, 2, 12kDa	AA195936	Hs.76362	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212350_at	0.016934	Consensus includes gb:AB029031.1 /DEF=Homo sapiens mRNA for KIAA1108 protein, partial cds. /FEA=mRNA /GEN=KIAA1108 /PROD=KIAA1108 protein /DB_XREF=gi:5689552 /UG=Hs.278586 KIAA1108 protein	AB029031		NP_055988
212355_at	1.26E-04	HCDI protein	AI075450	Hs.7911	
212356_at	0.00945	HCDI protein	AI075450	Hs.7911	
212367_at	0.016934	Consensus includes gb:AI799061 /FEA=EST /DB_XREF=gi:5364533 /DB_XREF=est:we98a10.x1 /CLONE=IMAGE:2349114 /UG=Hs.6048 FEM-1 (C.elegans) homolog b /FL=gb:AF178632.1 gb:NM_015322.1 gb:AF204883.1	NM_015322		NP_056137
212368_at	0.013223	hypothetical protein FLJ13564	AA972711	Hs.173081	
212369_at	3.26E-04	zinc finger protein 384	AI264312	Hs.103315	NP_597733
212370_x_at	0.041055	Consensus includes gb:AL080183.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O171 (from clone DKFZp434O171); partial cds. /FEA=mRNA /GEN=DKFZp434O171 /PROD=hypothetical protein /DB_XREF=gi:5262659 /UG=Hs.13273 KIAA0592 protein	AL080183		
212371_at	0.041055	Consensus includes gb:AL049397.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C1019 (from clone DKFZp586C1019). /FEA=mRNA /DB_XREF=gi:4500188 /UG=Hs.12314 Homo sapiens mRNA; cDNA DKFZp586C1019 (from clone DKFZp586C1019)	AL049397		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212373_at	0.001116	Consensus includes gb:AW139179 /FEA=EST /DB_XREF=gi:6143497 /DB_XREF=est:UI-H-BI1-aet-f-06-0-UI.s1 /CLONE=IMAGE:2720411 /UG=Hs.6048 FEM-1 (C.elegans) homolog b /FL=gb:AF178632.1 gb:NM_015322.1 gb:AF204883.1	NM_015322		NP_056137
212377_s_at	5.03E-04	Notch homolog 2 (Drosophila)	AU158495	Hs.131286	NP_077719
212382_at	0.029453	Consensus includes gb:BF433429 /FEA=EST /DB_XREF=gi:11445592 /DB_XREF=est:7q53a12.x1 /CLONE=IMAGE:3702046 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEMBB1000272	AK021980		
212383_at	0.005852	Consensus includes gb:AL096733.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E071 (from clone DKFZp434E071). /FEA=mRNA /DB_XREF=gi:5419866 /UG=Hs.267871 ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110116kD)	AL096733		NP_005168
212388_at	0.003862	Consensus includes gb:AB028980.1 /DEF=Homo sapiens mRNA for KIAA1057 protein, partial cds. /FEA=mRNA /GEN=KIAA1057 /PROD=KIAA1057 protein /DB_XREF=gi:5689450 /UG=Hs.7243 ubiquitin specific protease 24	AB028980		
212402_at	0.004341	KIAA0853 protein	BE895685	Hs.136102	NP_055885
212403_at	0.005852	Consensus includes gb:AI749193 /FEA=EST /DB_XREF=gi:5127457 /DB_XREF=est:at40e04.x1 /CLONE=IMAGE:2374494 /UG=Hs.17639 Homo sapiens ubiquitin protein ligase (UBE3B) mRNA, partial cds	AL096740		NP_569733

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212408_at	1.26E-04	Consensus includes gb:AK023204.1 /DEF=Homo sapiens cDNA FLJ13142 fis, clone NT2RP3003212, moderately similar to Rattus norvegicus lamina associated polypeptide 1C (LAP1C) mRNA. /FEA=mRNA /DB_XREF=gi:10435025 /UG=Hs.234265 DKFZP586G011 protein	AK023204		NP_056417
212412_at	0.003171	amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	AV715767	Hs.177486	
212421_at	0.00161	Consensus includes gb:AB023147.1 /DEF=Homo sapiens mRNA for KIAA0930 protein, partial cds. /FEA=mRNA /GEN=KIAA0930 /PROD=KIAA0930 protein /DB_XREF=gi:4589503 /UG=Hs.13255 KIAA0930 protein	AB023147		
212428_at	3.26E-04	Consensus includes gb:AW001101 /FEA=EST /DB_XREF=gi:5848017 /DB_XREF=est:wu24c05.x1 /CLONE=IMAGE:2520968 /UG=Hs.3852 KIAA0368 protein	AB002366		
212429_s_at	0.016686	general transcription factor IIIC, polypeptide 2, beta 110kDa	AW194657	Hs.75782	NP_001512
212433_x_at	0.005852	cadherin 1, type 1, E-cadherin (epithelial)	AA630314	Hs.356360	NP_002943
212434_at	0.016686	GrpE-like protein cochaperone	AL542571	Hs.151903	NP_079472
212436_at	7.44E-04	tripartite motif-containing 33	AI967961	Hs.287414	NP_148980
212437_at	0.033381	Consensus includes gb:AL109804 /DEF=Human DNA sequence from clone RP5-1009E24 on chromosome 20 Contains a novel gene encoding two isoforms similar to mouse sialoadhesin (a macrophage sialic acid binding receptor), a novel gene similar to KIAA0417, the CENPB gene (centromere protein ... /FEA=mRNA_8 /DB_XREF=gi:11121192 /UG=Hs.85004 centromere protein B (80kD)	AL109804		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212438_at	0.02145	putative nucleic acid binding protein RY-1	BG252325	Hs.54649	NP_006848
212440_at	0.013223	putative nucleic acid binding protein RY-1	BG252325	Hs.54649	NP_006848
212441_at	0.003171	Consensus includes gb:D86985.2 /DEF=Homo sapiens mRNA for KIAA0232 protein, partial cds. /FEA=mRNA /GEN=KIAA0232 /PROD=KIAA0232 protein /DB_XREF=gi:6634002 /UG=Hs.79276 KIAA0232 gene product	D86985		
212449_s_at	2.05E-04	lysophospholipase I	BG288007	Hs.409096	NP_006321
212451_at	0.018693	Consensus includes gb:N52532 /FEA=EST /DB_XREF=gi:1193698 /DB_XREF=est:yv35b11.s1 /CLONE=IMAGE:244701 /UG=Hs.118978 KIAA0256 gene product	D87445		
212452_x_at	8.68E-04	Consensus includes gb:AF113514.1 /DEF=Homo sapiens histone acetyltransferase MORF mRNA, complete cds. /FEA=CDS /PROD=histone acetyltransferase MORF /DB_XREF=gi:6002685 /UG=Hs.27590 histone acetyltransferase /FL=gb:AF113514.1	AF113514		NP_036462
212458_at	1.92E-04	ESTs, Weakly similar to SPY3_HUMAN Sprouty homolog 3 (Spry-3) [H.sapiens]	AW138902	Hs.392741	
212460_at	1.26E-04	hypothetical protein MGC24447	BE738425	Hs.353161	NP_612145
212461_at	1.26E-04	protease, serine, 15	BF793951	Hs.350265	NP_680479
212462_at	0.001618	Consensus includes gb:AU144267 /FEA=EST /DB_XREF=gi:11005788 /DB_XREF=est:AU144267 /CLONE=HEMBA1001396 /UG=Hs.27590 histone acetyltransferase /FL=gb:AF113514.1	AF113514		NP_036462
212463_at	0.004341	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	BE379006	Hs.42346	
212465_at	0.010205	hypothetical protein FLJ23027	AA524500	Hs.321766	NP_115609

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212467_at	1.26E-04	Consensus includes gb:AB014578.1 /DEF=Homo sapiens mRNA for KIAA0678 protein, partial cds. /FEA=mRNA /GEN=KIAA0678 /PROD=KIAA0678 protein /DB_XREF=gi:3327169 /UG=Hs.12707 KIAA0678 protein	AB014578		
212468_at	2.05E-04	Consensus includes gb:AK023512.1 /DEF=Homo sapiens cDNA FLJ13450 fis, clone PLACE1003027, highly similar to Homo sapiens mRNA for KIAA0516 protein. /FEA=mRNA /DB_XREF=gi:10435467 /UG=Hs.129872 sperm associated antigen 9	AB011088		NP_758853
212472_at	0.001116	KIAA0750 gene product	BE965029	Hs.314434	
212473_s_at	0.013223	KIAA0750 gene product	BE965029	Hs.314434	
212474_at	0.004341	KIAA0241 protein	BE503381	Hs.150275	
212476_at	0.00228	Consensus includes gb:D26069.1 /DEF=Human mRNA for KIAA0041 gene, partial cds. /FEA=mRNA /GEN=KIAA0041 /DB_XREF=gi:436227 /UG=Hs.24340 centaurin beta2	D26069		NP_036419
212477_at	0.001303	Consensus includes gb:D26069.1 /DEF=Human mRNA for KIAA0041 gene, partial cds. /FEA=mRNA /GEN=KIAA0041 /DB_XREF=gi:436227 /UG=Hs.24340 centaurin beta2	D26069		NP_036419
212479_s_at	0.007779	Consensus includes gb:AK022815.1 /DEF=Homo sapiens cDNA FLJ12753 fis, clone NT2RP2001226. /FEA=mRNA /DB_XREF=gi:10434432 /UG=Hs.75277 hypothetical protein FLJ13910	AL050139		NP_073617
212481_s_at	5.03E-04	tropomyosin 4	AI214061	Hs.250641	NP_003281

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212483_at	0.026891	Consensus includes gb:AW339587 /FEA=EST /DB_XREF=gi:6836213 /DB_XREF=est:he14g08.x1 /CLONE=IMAGE:2919038 /UG=Hs.225767 IDN3 protein	AB019494		NP_597677
212491_s_at	0.027243	DnaJ (Hsp40) homolog, subfamily C, member 8	AA843895	Hs.74711	NP_055095
212492_s_at	0.003171	KIAA0876 protein	AW237172	Hs.301011	
212496_s_at	0.00228	KIAA0876 protein	AW237172	Hs.301011	
212503_s_at	0.026891	KIAA0934 protein	N31807	Hs.227716	
212506_at	2.05E-04	ubiquitin specific protease 2	AL135735	Hs.199160	NP_009097
212507_at	1.26E-04	Consensus includes gb:D87446.1 /DEF=Human mRNA for KIAA0257 gene, partial cds. /FEA=mRNA /GEN=KIAA0257 /DB_XREF=gi:1665780 /UG=Hs.75912 KIAA0257 protein	D87446		
212511_at	0.003171	ubiquitin specific protease 2	AL135735	Hs.199160	NP_009097
212513_s_at	0.026891	Consensus includes gb:AB029020.1 /DEF=Homo sapiens mRNA for KIAA1097 protein, partial cds. /FEA=mRNA /GEN=KIAA1097 /PROD=KIAA1097 protein /DB_XREF=gi:5689530 /UG=Hs.173694 KIAA1097 protein	AB029020		NP_055832
212517_at	0.026891	Consensus includes gb:AL132773 /DEF=Human DNA sequence from clone RP4-741H3 on chromosome 20 Contains parts of isoform 1 and isoform 2 (KIAA0548) of the ATRN (attractin (with dipeptidylpeptidase IV activity)) gene, ESTs, STSs and GSSs /FEA=mRNA_1 /DB_XREF=gi:7159777 /UG=Hs.194019 attractin	AL132773		
212519_at	0.00161	ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast)	AL518159	Hs.7766	NP_003332

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212527_at	3.26E-04	Consensus includes gb:BF057059 /FEA=EST /DB_XREF=gi:10810955 /DB_XREF=est:7k15c07.x1 /CLONE=IMAGE:3444468 /UG=Hs.12421 hypothetical protein	AL023553		
212531_at	3.55E-04	Consensus includes gb:NM_005564.1 /DEF=Homo sapiens lipocalin 2 (oncogene 24p3) (LCN2), mRNA. /FEA=CDS /GEN=LCN2 /PROD=lipocalin 2 (oncogene 24p3) /DB_XREF=gi:5031852 /UG=Hs.204238 lipocalin 2 (oncogene 24p3) /FL=gb:NM_005564.1	NM_005564		NP_005555
212535_at	3.26E-04	MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	AA142929	Hs.277806	NP_005578
212536_at	1.26E-04	Consensus includes gb:AB023173.1 /DEF=Homo sapiens mRNA for KIAA0956 protein, partial cds. /FEA=mRNA /GEN=KIAA0956 /PROD=KIAA0956 protein /DB_XREF=gi:4589555 /UG=Hs.75478 ATPase, Class VI, type 11B	AB023173		
212542_s_at	0.041055	pleckstrin homology domain interacting protein	BF224151	Hs.367694	NP_060404
212543_at	1.26E-04	Consensus includes gb:U83115.1 /DEF=Human non-lens beta gamma-crystallin like protein (AIM1) mRNA, partial cds. /FEA=mRNA /GEN=AIM1 /PROD=non-lens beta gamma-crystallin like protein /DB_XREF=gi:2072424 /UG=Hs.161002 absent in melanoma 1	U83115		
212549_at	7.58E-04	Consensus includes gb:BE645861 /FEA=EST /DB_XREF=gi:9970172 /DB_XREF=est:7e77d05.x1 /CLONE=IMAGE:3288489 /UG=Hs.24064 Homo sapiens mRNA; cDNA DKFZp586N1323 (from clone DKFZp586N1323)	AL080218		
212552_at	0.001116	hippocalcin-like 1	BE617588	Hs.3618	NP_602293

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212553_at	0.033381	Consensus includes gb:AK026120.1 /DEF=Homo sapiens cDNA: FLJ22467 fis, clone HRC10345. /FEA=mRNA /DB_XREF=gi:10438867 /UG=Hs.29956 KIAA0460 protein	AK026120		
212560_at	0.016934	cytochrome c oxidase subunit VIb	AV728268	Hs.174031	
212561_at	0.003171	RAB6 interacting protein 1	AA349595	Hs.26797	
212570_at	0.009652	KIAA0830 protein	AL573201	Hs.167115	
212572_at	3.26E-04	Consensus includes gb:AW779556 /FEA=EST /DB_XREF=gi:7794159 /DB_XREF=est:hn81f05.x1 /CLONE=IMAGE:3034305 /UG=Hs.184523 KIAA0965 protein	AB023182		NP_055815
212583_at	0.011788	Consensus includes gb:AB011132.2 /DEF=Homo sapiens mRNA for KIAA0560 protein, partial cds. /FEA=mRNA /GEN=KIAA0560 /PROD=KIAA0560 protein /DB_XREF=gi:6635202 /UG=Hs.129952 KIAA0560 gene product	AB011132		
212584_at	0.010205	Consensus includes gb:BG260519 /FEA=EST /DB_XREF=gi:12770335 /DB_XREF=est:602372065F1 /CLONE=IMAGE:4480024 /UG=Hs.129952 KIAA0560 gene product	AB011132		
212586_at	0.02145	calpastatin	BG111635	Hs.9347	NP_775085
212590_at	7.58E-04	related RAS viral (r-ras) oncogene homolog 2	BG168858	Hs.206097	NP_036382
212594_at	0.033381	programmed cell death 4 (neoplastic transformation inhibitor)	N92498	Hs.326248	NP_663314
212597_s_at	1.26E-04	Consensus includes gb:AL079310.1 /DEF=Novel human gene mapping to chromosome 22. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:5102629 /UG=Hs.92260 high-mobility group protein 2-like 1	AL079310		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212599_at	0.00161	Consensus includes gb:AK025298.1 /DEF=Homo sapiens cDNA: FLJ21645 fis, clone COL08436. /FEA=mRNA /DB_XREF=gi:10437785 /UG=Hs.32168 KIAA0442 protein	AK025298		NP_056385
212601_at	0.004341	Consensus includes gb:AB007859.2 /DEF=Homo sapiens mRNA for KIAA0399 protein, partial cds. /FEA=mRNA /GEN=KIAA0399 /PROD=KIAA0399 protein /DB_XREF=gi:6634028 /UG=Hs.100955 KIAA0399 protein	AB007859		NP_055928
212602_at	2.05E-04	ALFY	AI806395	Hs.198135	NP_848700
212604_at	1.26E-04	Consensus includes gb:AI937794 /FEA=EST /DB_XREF=gi:5676664 /DB_XREF=est:wp82g10.x1 /CLONE=IMAGE:2468322 /UG=Hs.154655 imogen 38 /FL=gb:NM_005830.1	NM_005830		NP_005821
212606_at	1.26E-04	ALFY	AI806395	Hs.198135	NP_848700
212607_at	0.033381	Consensus includes gb:N32526 /FEA=EST /DB_XREF=gi:1152925 /DB_XREF=est:yy11f04.s1 /CLONE=IMAGE:270943 /UG=Hs.300642 serologically defined colon cancer antigen 8	U79271		NP_006633
212608_s_at	0.004341	ESTs, Moderately similar to Z254_HUMAN Zinc finger protein 254 (Bone marrow zinc finger 5) (BMZF-5) (Hematopoietic cell derived zinc finger protein 1) (HD-ZNF1) [H.sapiens]	W85912	Hs.409038	
212610_at	3.26E-04	Consensus includes gb:U79291.1 /DEF=Human clone 23721 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710271 /UG=Hs.83572 Human clone 23721 mRNA sequence	U79291		NP_116293

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212612_at	1.26E-04	Consensus includes gb:D31888.1 /DEF=Human mRNA for KIAA0071 gene, partial cds. /FEA=mRNA /GEN=KIAA0071 /DB_XREF=gi:506340 /UG=Hs.78398 KIAA0071 protein /FL=gb:AF155595.1 gb:NM_015156.1	D31888		NP_055971
212613_at	0.002418	butyrophilin, subfamily 3, member A2	AI991252	Hs.167741	NP_008978
212623_at	0.001116	Consensus includes gb:AU153138 /FEA=EST /DB_XREF=gi:11014659 /DB_XREF=est:AU153138 /CLONE=NT2RP3002507 /UG=Hs.174905 KIAA0033 protein	D26067		
212629_s_at	0.001054	Consensus includes gb:AI633689 /FEA=EST /DB_XREF=gi:4685019 /DB_XREF=est:th71d03.x1 /CLONE=IMAGE:2124101 /UG=Hs.69171 protein kinase C-like 2	AK023692		
212631_at	0.010205	Consensus includes gb:AI566082 /FEA=EST /DB_XREF=gi:4524534 /DB_XREF=est:tn53b12.x1 /CLONE=IMAGE:2172095 /UG=Hs.8906 Homo sapiens clone 24889 mRNA sequence	AF131808		NP_003560
212632_at	0.016934	Consensus includes gb:N32035 /FEA=EST /DB_XREF=gi:1152434 /DB_XREF=est:yw96e08.s1 /CLONE=IMAGE:260102 /UG=Hs.8906 Homo sapiens clone 24889 mRNA sequence	AF131808		NP_003560
212634_at	0.001618	KIAA0776 protein	AW298092	Hs.5460	NP_056138
212635_at	0.010205	ESTs	AW161626	Hs.409034	
212638_s_at	2.05E-04	VW domain-containing protein 1	BF131791	Hs.409154	NP_008944
212639_x_at	5.03E-04	tubulin, alpha, ubiquitous	AL581768	Hs.334842	NP_006073

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212642_s_at	0.041055	Consensus includes gb:AL023584 /DEF=Human DNA sequence from clone 67K17 on chromosome 6q24.1-24.3. Contains the HIVP2 (Schnurri-2) gene for HIV type 1 Enhancer-binding Protein 2, and a possible pseudogene in an intron of this gene. Contains STSs and GSSs and an AAAT repeat polymorp... /FEA=mRNA /DB_XREF=gi:3790154 /UG=Hs.75063 human immunodeficiency virus type 1 enhancer-binding protein 2 /FL=gb:NM_006734.1	AL023584		
212644_s_at	7.58E-04	likely ortholog of mouse MAPK-interacting and spindle-stabilizing protein	AI671747	Hs.388281	NP_653179
212649_at	0.004341	Consensus includes gb:AL079292.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 48814. /FEA=mRNA /PROD=hypothetical protein, similar to (AC007017)putative RNA helicase A Arabidopsis thaliana /DB_XREF=gi:5102732 /UG=Hs.95665 hypothetical protein	AL079292		NP_061903
212657_s_at	0.00161	interleukin 1 receptor antagonist	AW083357	Hs.81134	NP_776215
212658_at	0.00945	lipoma HMGIC fusion partner-like 2	N66633	Hs.79299	
212660_at	1.26E-04	KIAA0239 protein	AI735639	Hs.9729	NP_056103
212663_at	0.001116	Consensus includes gb:AB014574.1 /DEF=Homo sapiens mRNA for KIAA0674 protein, partial cds. /FEA=mRNA /GEN=KIAA0674 /PROD=KIAA0674 protein /DB_XREF=gi:3327161 /UG=Hs.14799 KIAA0674 protein	AB014574		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212666_at	7.58E-04	Consensus includes gb:AB046845.1 /DEF=Homo sapiens mRNA for KIAA1625 protein, partial cds. /FEA=mRNA /GEN=KIAA1625 /PROD=KIAA1625 protein /DB_XREF=gi:10047326 /UG=Hs.119120 E3 ubiquitin ligase SMURF1	AB046845		NP_851994
212669_at	0.003171	calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	AI093569	Hs.153406	NP_751913
212671_s_at	0.003171	major histocompatibility complex, class II, DQ alpha 1	BG397856	Hs.198253	NP_002113
212674_s_at	5.03E-04	Consensus includes gb:AK002076.1 /DEF=Homo sapiens cDNA FLJ11214 fis, clone PLACE1007990. /FEA=mRNA /DB_XREF=gi:7023738 /UG=Hs.281616 Homo sapiens cDNA FLJ11214 fis, clone PLACE1007990	AK002076		NP_619520
212677_s_at	0.004341	Consensus includes gb:BG530481 /FEA=EST /DB_XREF=gi:13522018 /DB_XREF=est:602559046F1 /CLONE=IMAGE:4697218 /UG=Hs.79507 KIAA0582 protein	AB011154		NP_055962
212678_at	0.011377	neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	AW293356	Hs.93207	
212683_at	3.56E-04	KIAA0446 gene product	AL526243	Hs.158286	
212685_s_at	3.26E-04	Consensus includes gb:AI608789 /FEA=EST /DB_XREF=gi:4617956 /DB_XREF=est:tw94g12.x1 /CLONE=IMAGE:2267398 /UG=Hs.52515 transducin (beta)-like 2	AK026529		NP_116783

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212687_at	0.013223	Consensus includes gb:AL110164.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586l0324 (from clone DKFZp586l0324). /FEA=mRNA /DB_XREF=gi:5817069 /UG=Hs.193700 Homo sapiens mRNA; cDNA DKFZp586l0324 (from clone DKFZp586l0324)	AL110164		
212688_at	1.26E-04	Consensus includes gb:BC003393.1 /DEF=Homo sapiens, clone IMAGE:3447173, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3447173) /DB_XREF=gi:13097275 /UG=Hs.239818 phosphoinositide-3-kinase, catalytic, beta polypeptide /FL=gb:NM_006219.1	BC003393		NP_006210
212689_s_at	0.007779	Human putative zinc finger protein mRNA	AA524505	Hs.383008	NP_060903
212697_at	0.010205	Lutheran blood group (Auberger b antigen included)	AL515874	Hs.155048	NP_835227
212698_s_at	0.02008	hypothetical protein FLJ11619	BF966021	Hs.79844	NP_848699
212702_s_at	1.26E-04	Consensus includes gb:N45111 /FEA=EST /DB_XREF=gi:1186277 /DB_XREF=est:yz12f12.s1 /CLONE=IMAGE:282863 /UG=Hs.330988 Homo sapiens, Similar to Bicaudal D (Drosophila) homolog 1, clone IMAGE:3622452, mRNA, partial cds	BC004296		NP_056065
212710_at	0.025245	DKFZP434F195 protein	BE222901	Hs.184793	
212717_at	5.03E-04	Consensus includes gb:AB002354.2 /DEF=Homo sapiens mRNA for KIAA0356 protein, partial cds. /FEA=mRNA /GEN=KIAA0356 /PROD=KIAA0356 protein /DB_XREF=gi:6634022 /UG=Hs.32312 KIAA0356 gene product	AJ002220		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212722_s_at	0.026891	Consensus includes gb:AK021780.1 /DEF=Homo sapiens cDNA FLJ11718 fis, clone HEMBA1005252, highly similar to Homo sapiens mRNA for KIAA0585 protein. /FEA=mRNA /DB_XREF=gi:10433034 /UG=Hs.72660 phosphatidylserine receptor	AK021780		NP_055982
212723_at	1.26E-04	Consensus includes gb:AK021780.1 /DEF=Homo sapiens cDNA FLJ11718 fis, clone HEMBA1005252, highly similar to Homo sapiens mRNA for KIAA0585 protein. /FEA=mRNA /DB_XREF=gi:10433034 /UG=Hs.72660 phosphatidylserine receptor	AK021780		NP_055982
212725_s_at	0.026891	ESTs	N37081	Hs.409222	
212726_at	0.00161	Consensus includes gb:AB014562.1 /DEF=Homo sapiens mRNA for KIAA0662 protein, partial cds. /FEA=mRNA /GEN=KIAA0662 /PROD=KIAA0662 protein /DB_XREF=gi:3327137 /UG=Hs.93868 KIAA0662 gene product	AB014562		NP_078793
212737_at	0.00161	GM2 ganglioside activator protein	AL513583	Hs.289082	NP_000396
212740_at	0.001116	phosphoinositide-3-kinase, regulatory subunit 4, p150	BF740111	Hs.83050	NP_055417
212742_at	1.26E-04	zinc finger protein 364	AL530462	Hs.16537	
212743_at	0.007781	Consensus includes gb:AV694093 /FEA=EST /DB_XREF=gi:10295956 /DB_XREF=est:AV694093 /CLONE=GKCDSD10 /UG=Hs.48297 DKFZP586C1620 protein	AL050144		NP_056251
212744_at	0.002671	Bardet-Biedl syndrome 4	AI813772	Hs.387416	NP_149017
212747_at	1.28E-04	KIAA0229 protein	AI990523	Hs.221332	
212749_s_at	1.26E-04	Consensus includes gb:AI096477 /FEA=EST /DB_XREF=gi:3445971 /DB_XREF=est:qa03c06.x1 /CLONE=IMAGE:1685674 /UG=Hs.48297 DKFZP586C1620 protein	AL050144		NP_056251

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212752_at	7.58E-04	cytoplasmic linker associated protein 1	AA176798	Hs.392661	
212757_s_at	0.033381	calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	BF111268	Hs.153406	NP_751913
212761_at	0.016074	transcription factor 7-like 2 (T-cell specific, HMG-box)	AI703074	Hs.348412	NP_110383
212768_s_at	7.31E-04	isoform 1 match: proteins: Sw:Q07081 Tr:O95362 Tr:Q9Z2Y4 Tr:O95897 Tr:O70624 Sw:Q99972 Sw:Q99784 Sw:Q62609 Tr:Q9TV76 Tr:Q9I9K5 Sw:P01813 Tr:Q9IAK4 Tr:O35429; Human DNA sequence from clone RP11-209J19 on chromosome 13 Contains ESTs, STSs and GSSs. Contains the gene for the GW112 protein with two isoforms (GW112 and KIAA4294), complete sequence.	AL390736		
212769_at	0.026891	transducin-like enhancer of split 3 (E(sp1) homolog, Drosophila)	AI567426	Hs.156183	NP_005069
212779_at	0.004341	Consensus includes gb:AB029032.1 /DEF=Homo sapiens mRNA for KIAA1109 protein, partial cds. /FEA=mRNA /GEN=KIAA1109 /PROD=KIAA1109 protein /DB_XREF=gi:5689554 /UG=Hs.6606 KIAA1109 protein	AB029032		NP_115578
212780_at	0.007779	Consensus includes gb:AA700167 /FEA=EST /DB_XREF=gi:2703130 /DB_XREF=est:zj36h12.s1 /CLONE=IMAGE:452423 /UG=Hs.326392 son of sevenless (Drosophila) homolog 1 /FL=gb:L13857.1	L13857		NP_005624
212783_at	1.26E-04	Consensus includes gb:AI538172 /FEA=EST /DB_XREF=gi:4452307 /DB_XREF=est:ti75f08.x1 /CLONE=IMAGE:2137863 /UG=Hs.91065 hypothetical protein DKFZp761B2423	AK026954		NP_008841
212786_at	0.016934	KIAA0350 protein	AA731693	Hs.23263	
212787_at	3.26E-04	ZAP3 protein	AI952986	Hs.159471	
212790_x_at	0.010205	ribosomal protein L13a	BF942308	Hs.119122	NP_775866

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212794_s_at	0.013223	Consensus includes gb:AK001728.1 /DEF=Homo sapiens cDNA FLJ10866 fis, clone NT2RP4001614. /FEA=mRNA /DB_XREF=gi:7023169 /UG=Hs.12144 KIAA1033 protein	AL137753		
212796_s_at	0.041055	KIAA1055 protein	BF195608	Hs.126084	NP_055894
212799_at	3.26E-04	Consensus includes gb:BE217875 /FEA=EST /DB_XREF=gi:8905193 /DB_XREF=est:hv31a03.x1 /CLONE=IMAGE:3174988 /UG=Hs.12311 Homo sapiens clone 23570 mRNA sequence	AF038202		NP_005810
212802_s_at	7.58E-04	Consensus includes gb:AK023841.1 /DEF=Homo sapiens cDNA FLJ13779 fis, clone PLACE4000445, highly similar to Homo sapiens mRNA; cDNA DKFZp434C212 (from clone DKFZp434C212). /FEA=mRNA /DB_XREF=gi:10435900 /UG=Hs.172069 DKFZP434C212 protein	AK023841		
212804_s_at	0.026891	Consensus includes gb:AI797397 /FEA=EST /DB_XREF=gi:5362869 /DB_XREF=est:we87f12.x1 /CLONE=IMAGE:2348111 /UG=Hs.172069 DKFZP434C212 protein	AK023841		
212807_s_at	3.26E-04	sortilin 1	BE742268	Hs.351872	NP_002950
212808_at	0.041055	hypothetical protein FLJ14639	AA152202	Hs.334788	NP_116204
212811_x_at	5.03E-04	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	BG032165	Hs.323878	NP_003029
212814_at	0.033381	Consensus includes gb:AB020635.1 /DEF=Homo sapiens mRNA for KIAA0828 protein, partial cds. /FEA=mRNA /GEN=KIAA0828 /PROD=KIAA0828 protein /DB_XREF=gi:4240144 /UG=Hs.4984 KIAA0828 protein	AB020635		NP_056143
212823_s_at	0.007052	KIAA0599 protein	AU147160	Hs.198037	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212827_at	1.26E-04	Consensus includes gb:X17115.1 /DEF=Human mRNA for IgM heavy chain complete sequence. /FEA=mRNA /DB_XREF=gi:33450 /UG=Hs.302063 immunoglobulin heavy constant mu	X17115		
212828_at	0.009199	Consensus includes gb:AA191573 /FEA=EST /DB_XREF=gi:1780272 /DB_XREF=est:zp81b09.s1 /CLONE=IMAGE:626585 /UG=Hs.61289 synaptotagmin 2	AL157424		NP_003889
212829_at	0.010205	ESTs	BE878277	Hs.57079	
212830_at	2.05E-04	EGF-like-domain, multiple 5	BF110421	Hs.5599	
212833_at	0.007779	Consensus includes gb:M74089.1 /DEF=Human TB1 gene mRNA, 3 end. /FEA=mRNA /GEN=TB1 /DB_XREF=gi:182400 /UG=Hs.75639 Human TB1 gene mRNA, 3 end	M74089		NP_620128
212836_at	0.016934	Consensus includes gb:D26018.1 /DEF=Human mRNA for KIAA0039 gene, partial cds. /FEA=mRNA /GEN=KIAA0039 /DB_XREF=gi:436221 /UG=Hs.82502 polymerase (DNA directed), delta 3	D26018		
212838_at	0.004452	Consensus includes gb:AB023227.1 /DEF=Homo sapiens mRNA for KIAA1010 protein, partial cds. /FEA=mRNA /GEN=KIAA1010 /PROD=KIAA1010 protein /DB_XREF=gi:4589669 /UG=Hs.23860 KIAA1010 protein	AB023227		
212840_at	0.016934	Consensus includes gb:BG339560 /FEA=EST /DB_XREF=gi:13145998 /DB_XREF=est:602437413F1 /CLONE=IMAGE:4555466 /UG=Hs.127287 KIAA0794 protein	AB018337		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212854_x_at	0.003171	Consensus includes gb:AB051480.1 /DEF=Homo sapiens mRNA for KIAA1693 protein, partial cds. /FEA=mRNA /GEN=KIAA1693 /PROD=KIAA1693 protein /DB_XREF=gi:12697930 /UG=Hs.323463 KIAA1693 protein; hypothetical protein FLJ20719	AB051480		NP_060410
212856_at	1.26E-04	Consensus includes gb:AB018310.1 /DEF=Homo sapiens mRNA for KIAA0767 protein, partial cds. /FEA=mRNA /GEN=KIAA0767 /PROD=KIAA0767 protein /DB_XREF=gi:3882254 /UG=Hs.95180 KIAA0767 protein	AB018310		
212861_at	0.003171	hypothetical protein MGC11308	BF690150	Hs.19210	NP_116278
212862_at	0.016934	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2	AL568982	Hs.24812	NP_003809
212864_at	0.00161	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2	AL568982	Hs.24812	NP_003809
212867_at	0.001116	nuclear receptor coactivator 2	AI040324	Hs.29131	NP_006531
212869_x_at	0.026891	tumor protein, translationally-controlled 1	AI721229	Hs.279860	NP_003286
212870_at	1.26E-04	son of sevenless homolog 2 (Drosophila)	AI628605	Hs.348496	
212871_at	0.020576	Consensus includes gb:NM_003668.1 /DEF=Homo sapiens mitogen-activated protein kinase-activated protein kinase 5 (MAPKAPK5), mRNA. /FEA=CDS /GEN=MAPKAPK5 /PROD=mitogen-activated protein kinase-activated protein kinase 5 /DB_XREF=gi:4506034 /UG=Hs.30327 mitogen-activated protein kinase-activated protein kinase 5 /FL=gb:NM_003668.1	NM_003668		NP_620777

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212872_s_at	0.005852	Consensus includes gb:AK023092.1 /DEF=Homo sapiens cDNA FLJ13030 fis, clone NT2RP3001111, highly similar to Homo sapiens TRF-proximal protein mRNA. /FEA=mRNA /DB_XREF=gi:10434851 /UG=Hs.93796 DKFZP586D2223 protein	AK023092		NP_061031
212880_at	0.041055	Consensus includes gb:AB011113.1 /DEF=Homo sapiens mRNA for KIAA0541 protein, partial cds. /FEA=mRNA /GEN=KIAA0541 /PROD=KIAA0541 protein /DB_XREF=gi:3043605 /UG=Hs.10881 WD repeat domain 7	AB011113		NP_443066
212882_at	3.26E-04	Consensus includes gb:AB018338.1 /DEF=Homo sapiens mRNA for KIAA0795 protein, partial cds. /FEA=mRNA /GEN=KIAA0795 /PROD=KIAA0795 protein /DB_XREF=gi:3882310 /UG=Hs.22926 KIAA0795 protein	AB018338		NP_079286
212887_at	1.26E-04	Sec23 homolog A (S. cerevisiae)	AI753659	Hs.272927	NP_006355
212888_at	0.007779	ESTs, Highly similar to A32915 nucleophosmin - human [H.sapiens]	BG109746	Hs.409325	
212890_at	0.003171	hypothetical protein MGC15523	BF663461	Hs.381200	NP_612637
212895_s_at	7.58E-04	active BCR-related gene	AL527773	Hs.408408	NP_068781
212899_at	7.58E-04	Consensus includes gb:AB028951.1 /DEF=Homo sapiens mRNA for KIAA1028 protein, partial cds. /FEA=mRNA /GEN=KIAA1028 /PROD=KIAA1028 protein /DB_XREF=gi:5689392 /UG=Hs.129836 KIAA1028 protein	AL122055		NP_055891

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212908_at	0.037567	Consensus includes gb:AK022530.1 /DEF=Homo sapiens cDNA FLJ12468 fis, clone NT2RM1000857, highly similar to Homo sapiens mRNA for KIAA0962 protein. /FEA=mRNA /DB_XREF=gi:10433971 /UG=Hs.9059 KIAA0962 protein	AB023179		
212911_at	0.041055	Consensus includes gb:AB023179.1 /DEF=Homo sapiens mRNA for KIAA0962 protein, partial cds. /FEA=mRNA /GEN=KIAA0962 /PROD=KIAA0962 protein /DB_XREF=gi:4589567 /UG=Hs.9059 KIAA0962 protein	AB023179		
212916_at	5.03E-04	KIAA1111 protein	AW249934	Hs.285641	
212917_x_at	0.003171	RecQ protein-like (DNA helicase Q1-like)	BF219234	Hs.235069	NP_079130
212920_at	0.010205	ESTs	AV682285	Hs.227630	
212926_at	0.008036	Consensus includes gb:AW183677 /FEA=EST /DB_XREF=gi:6452191 /DB_XREF=est:xj86e07.x1 /CLONE=IMAGE:2664132 /UG=Hs.103283 KIAA0594 protein	AB011166		NP_055925
212929_s_at	0.016686	hypothetical protein FLJ10824	W68158	Hs.13273	
212931_at	0.00228	Consensus includes gb:AB006630.1 /DEF=Homo sapiens mRNA for KIAA0292 gene, partial cds. /FEA=mRNA /GEN=KIAA0292 /DB_XREF=gi:2564331 /UG=Hs.201668 transcription factor 20 (AR1)	AB006630		
212932_at	0.013088	Consensus includes gb:AK022494.1 /DEF=Homo sapiens cDNA FLJ12432 fis, clone NT2RM1000018, highly similar to Human mRNA for KIAA0066 gene. /FEA=mRNA /DB_XREF=gi:10433912 /UG=Hs.227881 RAB3 GTPase-ACTIVATING PROTEIN	AK022494		
212934_at	1.26E-04	ESTs, Weakly similar to p47 [Homo sapiens] [H.sapiens]	AI245523	Hs.131357	
212936_at	0.013223	hypothetical protein DKFZp564D172	AI927701	Hs.25307	NP_114431

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212947_at	0.007779	Consensus includes gb:AL031685 /DEF=Human DNA sequence from clone RP5-963K23 on chromosome 20q13.11-13.2 Contains a KRT18 (Keratin type I, Cytoskeletal 18 (Cytokeratin 18, CK18,CYK18)) pseudogene, a gene for a novel protein, the gene for spermatogenesis associated protein PD1 (KIAA0... /FEA=mRNA_3 /DB_XREF=gi:9368423 /UG=Hs.12785 KIAA0939 protein	AL031685		
212952_at	0.042466	calreticulin	BE251303	Hs.16488	NP_004334
212953_x_at	0.009907	calreticulin	BE251303	Hs.16488	NP_004334
212958_x_at	0.003171	peptidylglycine alpha-amidating monooxygenase	AI022882	Hs.83920	NP_620177
212961_x_at	6.68E-04	Consensus includes gb:L43578.1 /DEF=Homo sapiens (clone 115392) mRNA. /FEA=mRNA /DB_XREF=gi:899067 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43578		NP_835225
212971_at	0.007779	cysteinyl-tRNA synthetase	AI769685	Hs.159604	NP_644802
212973_at	3.26E-04	ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase)	AI692341	Hs.79886	NP_653164
212975_at	0.005852	Consensus includes gb:AB020677.2 /DEF=Homo sapiens mRNA for KIAA0870 protein, partial cds. /FEA=mRNA /GEN=KIAA0870 /PROD=KIAA0870 protein /DB_XREF=gi:6635136 /UG=Hs.18166 KIAA0870 protein	AB020677		
212982_at	3.26E-04	Huntingtin interacting protein H	AI621223	Hs.4014	NP_056151
212984_at	0.026891	activating transcription factor 2	BE786164	Hs.198166	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212987_at	1.26E-04	Consensus includes gb:AL031178 /DEF=Human DNA sequence from clone 341E18 on chromosome 6p11.2-12.3. Contains a SerineThreonine Protein Kinase gene (presumptive isolog of a Rat gene) and a novel alternatively spliced gene. Contains a putative CpG island, ESTs and GSSs /FEA=mRNA_2 /DB_XREF=gi:3550040 /UG=Hs.11050 F-box only protein 9	AL137520		NP_258442
212990_at	1.26E-04	Consensus includes gb:AB020717.1 /DEF=Homo sapiens mRNA for KIAA0910 protein, partial cds. /FEA=mRNA /GEN=KIAA0910 /PROD=KIAA0910 protein /DB_XREF=gi:4240308 /UG=Hs.127416 synaptojanin 1	AB020717		NP_003886
212991_at	0.003171	Consensus includes gb:AL137520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C0118 (from clone DKFZp434C0118); partial cds. /FEA=mRNA /GEN=DKFZp434C0118 /PROD=hypothetical protein /DB_XREF=gi:6808183 /UG=Hs.11050 F-box only protein 9	AL137520		NP_258442
212993_at	0.004341	ESTs, Weakly similar to hypothetical protein FLJ12547 [Homo sapiens] [H.sapiens]	AA114166	Hs.381190	
213006_at	0.013223	CCAAT/enhancer binding protein (C/EBP), delta	AV655640	Hs.76722	NP_005186
213009_s_at	0.035171	Consensus includes gb:AK022701.1 /DEF=Homo sapiens cDNA FLJ12639 fis, clone NT2RM4001938, highly similar to Homo sapiens mRNA for KIAA0898 protein. /FEA=mRNA /DB_XREF=gi:10434250 /UG=Hs.8164 Mulibrey nanism	AK022701		NP_056109
213011_s_at	2.05E-04	triosephosphate isomerase 1	BF116254	Hs.83848	NP_000356

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213012_at	1.28E-04	Consensus includes gb:D42055.1 /DEF=Human mRNA for KIAA0093 gene, partial cds. /FEA=mRNA /GEN=KIAA0093 /DB_XREF=gi:577312 /UG=Hs.1565 neural precursor cell expressed, developmentally down-regulated 4	D42055		
213014_at	0.041795	Consensus includes gb:BG222394 /FEA=EST /DB_XREF=gi:12707915 /DB_XREF=est:naf75h09.x1 /CLONE=IMAGE:4170089 /UG=Hs.234249 mitogen-activated protein kinase 8 interacting protein 1 /FL=gb:AF074091.1 gb:NM_005456.1	NM_005456		NP_005447
213015_at	0.003539	ESTs	BF448315	Hs.171553	
213016_at	1.26E-04	ESTs	BF448315	Hs.171553	
213022_s_at	0.041055	Consensus includes gb:NM_007124.1 /DEF=Homo sapiens utrophin (homologous to dystrophin) (UTRN), mRNA. /FEA=CDS /GEN=UTRN /PROD=utrophin /DB_XREF=gi:6005937 /UG=Hs.251967 utrophin (homologous to dystrophin) /FL=gb:NM_007124.1	NM_007124		NP_009055
213027_at	0.007779	Sjogren syndrome antigen A2 (60kDa, ribonucleoprotein autoantigen SS-A/Ro)	AU146655	Hs.554	
213037_x_at	5.03E-04	Consensus includes gb:AJ132258.1 /DEF=Homo sapiens mRNA for staufen protein, partial. /FEA=mRNA /GEN=staufen /PROD=staufen protein /DB_XREF=gi:4572587 /UG=Hs.6113 staufen (Drosophila, RNA-binding protein)	AJ132258		NP_059348
213044_at	0.001116	Rho-associated, coiled-coil containing protein kinase 1	N22548	Hs.17820	NP_005397
213048_s_at	1.26E-04	SET translocation (myeloid leukemia-associated)	W26593	Hs.145279	
213049_at	0.040064	hypothetical protein DKFZp667F074	BG436400	Hs.378933	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213052_at	0.041055	protein kinase, cAMP-dependent, regulatory, type II, alpha	BF246917	Hs.365523	NP_004148
213056_at	1.26E-04	KIAA1013 protein	AU145019	Hs.96427	
213061_s_at	0.00161	Homo sapiens, Similar to N-terminal Asn amidase, clone MGC:29626 IMAGE:4872717, mRNA, complete cds	AA643304	Hs.351573	NP_775745
213073_at	0.028067	Consensus includes gb:AB002319.1 /DEF=Human mRNA for KIAA0321 gene, partial cds. /FEA=mRNA /GEN=KIAA0321 /DB_XREF=gi:2224582 /UG=Hs.8663 KIAA0321 protein	AB002319		
213084_x_at	0.00161	cadherin 1, type 1, E-cadherin (epithelial)	BF125158	Hs.356360	NP_000975
213086_s_at	0.016934	casein kinase 1, alpha 1	BF341845	Hs.283738	NP_001883
213095_x_at	5.03E-04	Consensus includes gb:AF299327.1 /DEF=Homo sapiens allograft inflammatory factor-1 splice variant G1 mRNA, complete cds. /FEA=CDS /PROD=allograft inflammatory factor-1 splice variantG1 /DB_XREF=gi:11611961 /UG=Hs.76364 allograft inflammatory factor 1 /FL=gb:AF299327.1	AF299327		NP_116573
213102_at	0.005852	ARP3 actin-related protein 3 homolog (yeast)	Z78330	Hs.5321	NP_005712
213109_at	0.009689	KIAA0551 protein	N25621	Hs.170204	
213111_at	0.02145	Consensus includes gb:AB023198.1 /DEF=Homo sapiens mRNA for KIAA0981 protein, partial cds. /FEA=mRNA /GEN=KIAA0981 /PROD=KIAA0981 protein /DB_XREF=gi:4589605 /UG=Hs.158135 KIAA0981 protein	AB023198		
213113_s_at	0.005852	likely ortholog of mouse embryonic epithelial gene 1	AI630178	Hs.274453	NP_060081

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213115_at	0.005852	Consensus includes gb:AL031177 /DEF=Human DNA sequence from clone 889N15 on chromosome Xq22.1-22.3. Contains part of the gene for a novel protein similar to X. laevis Cortical Thymocyte Marker CTX, the possibly alternatively spliced gene for 26S Proteasome subunit p28 (Ankyrin repea... /FEA=mRNA_3 /DB_XREF=gi:4071056 /UG=Hs.8763 Human DNA sequence from clone 889N15 on chromosome Xq22.1-22.3. Contains part of the gene for a novel protein similar to X. laevis Cortical Thymocyte Marker CTX, the possibly alternatively spliced gene for 26S Proteasome subunit p28 (Ankyrin repeat protei	AL031177		
213117_at	5.03E-04	KIAA1354 protein	AW138594	Hs.106283	NP_061335
213118_at	2.17E-04	Consensus includes gb:AL136821.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M1526 (from clone DKFZp434M1526). /FEA=mRNA /GEN=DKFZp434M1526 /PROD=hypothetical protein /DB_XREF=gi:12053152 /UG=Hs.153293 KIAA0701 protein	AL136821		
213119_at	0.003368	ESTs	AW058600	Hs.392501	
213120_at	0.009582	Consensus includes gb:AA405798 /FEA=EST /DB_XREF=gi:2063781 /DB_XREF=est:zu57f04.s1 /CLONE=IMAGE:742111 /UG=Hs.153293 KIAA0701 protein	AL136821		
213123_at	1.26E-04	cytochrome c oxidase subunit Vb	BE222709	Hs.1342	
213127_s_at	0.001911	mediator of RNA polymerase II transcription subunit MED8	BG230758	Hs.301756	NP_443109
213128_s_at	0.001116	ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome)	AA527499	Hs.180686	NP_570854

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213133_s_at	0.001911	ESTs, Highly similar to GCHUH glycine cleavage system protein H precursor - human [H.sapiens]	BE908931	Hs.356054	
213137_s_at	0.042536	protein tyrosine phosphatase, non-receptor type 2	AI828880	Hs.82829	NP_536348
213145_at	0.041055	hypothetical protein MGC40195	BF001666	Hs.367956	
213154_s_at	2.05E-04	Consensus includes gb:AI934125 /FEA=EST /DB_XREF=gi:5672995 /DB_XREF=est:wn97c08.x1 /CLONE=IMAGE:2453774 /UG=Hs.17411 KIAA0699 protein	AB014599		NP_056065
213158_at	0.016934	Consensus includes gb:AA045174 /FEA=EST /DB_XREF=gi:1523376 /DB_XREF=est:zk66a07.s1 /CLONE=IMAGE:487764 /UG=Hs.16193 Homo sapiens mRNA; cDNA DKFZp586B211 (from clone DKFZp586B211)	AL049423		
213159_at	2.17E-04	Consensus includes gb:AB018348.1 /DEF=Homo sapiens mRNA for KIAA0805 protein, partial cds. /FEA=mRNA /GEN=KIAA0805 /PROD=KIAA0805 protein /DB_XREF=gi:3882330 /UG=Hs.55947 KIAA0805 protein	AB018348		NP_055797
213160_at	1.26E-04	Consensus includes gb:D86964.1 /DEF=Human mRNA for KIAA0209 gene, partial cds. /FEA=mRNA /GEN=KIAA0209 /DB_XREF=gi:1504001 /UG=Hs.17211 dedicator of cytokinesis 2	D86964		NP_004937
213165_at	0.00228	centrosome-associated protein 350	AI041204	Hs.92200	
213166_x_at	0.030771	Homo sapiens, clone IMAGE:4634703, mRNA, partial cds	BG332462	Hs.355815	
213173_at	1.49E-04	Consensus includes gb:AI815033 /FEA=EST /DB_XREF=gi:5426248 /DB_XREF=est:wk70c01.x1 /CLONE=IMAGE:2420736 /UG=Hs.55947 KIAA0805 protein	AB018348		NP_055797

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213193_x_at	0.00161	T cell receptor beta locus	AL559122	Hs.155664	
213198_at	1.26E-04	Consensus includes gb:AL117643.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M245 (from clone DKFZp434M245). /FEA=mRNA /DB_XREF=gi:5912233 /UG=Hs.5288 Homo sapiens mRNA; cDNA DKFZp434M245 (from clone DKFZp434M245)	AL117643		NP_064733
213208_at	0.007779	KIAA0240 protein	AI801951	Hs.196275	
213215_at	0.00228	Homo sapiens, clone IMAGE:4157286, mRNA, partial cds	AI910895	Hs.392619	
213216_at	0.031704	KIAA0459 protein	AL537463	Hs.374987	
213220_at	0.033381	six transmembrane epithelial antigen of prostate 2	AV706096	Hs.409158	
213222_at	0.040938	continued from dJ881E24.1.1 in Em:AL031683 match: proteins: Sw:P10687 Tr:Q9Z1B3 Sw:P10894; Human DNA sequence from clone RP4-654A7 on chromosome 20 Contains the 3' end of two variants of the PLCB1 gene for phospholipase C beta 1 (phosphoinositide-specific), complete sequence.	AL049593		
213225_at	1.26E-04	Consensus includes gb:AJ271832.1 /DEF=Homo sapiens mRNA for protein phosphatase 1B2 (PPM1B2 gene). /FEA=mRNA /GEN=PPM1B2 /PROD=protein phosphatase 1B2 53 kDa isoform /DB_XREF=gi:12666516 /UG=Hs.5687 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform /FL=gb:NM_002706.1	AJ271832		NP_808908
213229_at	2.05E-04	Dicer1, Dcr-1 homolog (Drosophila)	BF590131	Hs.87889	NP_803187
213241_at	1.26E-04	Consensus includes gb:AF035307.1 /DEF=Homo sapiens clone 23785 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661068 /UG=Hs.184697 Homo sapiens clone 23785 mRNA sequence	AF035307		
213243_at	0.016934	KIAA0532 protein	AI052003	Hs.21077	
213246_at	1.26E-04	DKFZP564F1123 protein	AI346504	Hs.275352	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213254_at	5.03E-04	KIAA1093 protein	N64803	Hs.117333	
213279_at	0.00161	Consensus includes gb:AL050217.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586l0523 (from clone DKFZp586l0523). /FEA=mRNA /DB_XREF=gi:4884458 /UG=Hs.308467 Homo sapiens mRNA; cDNA DKFZp586l0523 (from clone DKFZp586l0523)	AL050217		
213280_at	0.026891	Consensus includes gb:AK000478.1 /DEF=Homo sapiens cDNA FLJ20471 fis, clone KAT06974. /FEA=mRNA /DB_XREF=gi:7020593 /UG=Hs.301552 KIAA1039 protein	AK000478		
213288_at	1.41E-04	ESTs	AI761250	Hs.396081	
213292_s_at	0.001116	sorting nexin 13	AA908770	Hs.283881	NP_055947
213296_at	0.040938	peroxisome biogenesis factor 10	BF339133	Hs.247220	NP_722540
213301_x_at	0.001911	transcriptional intermediary factor 1	AL538264	Hs.183858	NP_003843
213305_s_at	0.013223	Consensus includes gb:L42375.1 /DEF=Homo sapiens protein phosphatase 2A B56-gamma1 (PP2A) mRNA, 3 end of cds. /FEA=mRNA /GEN=PP2A /PROD=protein phosphatase 2A B56-gamma1 /DB_XREF=gi:1000891 /UG=Hs.171734 protein phosphatase 2, regulatory subunit B (B56), gamma isoform	L42375		NP_848703
213309_at	0.004341	Consensus includes gb:AL117515.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L0735 (from clone DKFZp434L0735); partial cds. /FEA=mRNA /GEN=DKFZp434L0735 /PROD=hypothetical protein /DB_XREF=gi:5912029 /UG=Hs.54886 phospholipase C, epsilon 2	AL117515		NP_055999
213311_s_at	0.003539	KIAA1049 protein	BF000251	Hs.227835	NP_055787
213313_at	0.001116	rab6 GTPase activating protein (GAP and centrosome-associated)	AI922519	Hs.55099	NP_036329

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213314_at	0.027633	Consensus includes gb:NM_020425.1 /DEF=Homo sapiens hypothetical protein DKFZp586E1923 (DKFZP586E1923), mRNA. /FEA=CDS /GEN=DKFZP586E1923 /PROD=hypothetical protein DKFZp586E1923 /DB_XREF=gi:10092684 /UG=Hs.70769 hypothetical protein DKFZp586E1923 /FL=gb:NM_020425.1	NM_020425		NP_065158
213315_x_at	0.005411	Homo sapiens, clone MGC:10039 IMAGE:3889701, mRNA, complete cds	L43577	Hs.82171	NP_835225
213318_s_at	0.041055	apolipoprotein M	BG028844	Hs.274348	NP_542434
213327_s_at	0.004341	ubiquitin specific protease 12	AI820101	Hs.42400	
213341_at	0.004631	fem-1 homolog c (C.elegans)	AI862658	Hs.47367	NP_064562
213344_s_at	0.003171	H2A histone family, member X	H51429	Hs.147097	NP_002096
213351_s_at	0.008185	KIAA0779 protein	AI934469	Hs.362996	
213355_at	0.014715	Consensus includes gb:AI989567 /FEA=EST /DB_XREF=gi:5836448 /DB_XREF=est:ws34e03.x1 /CLONE=IMAGE:2499100 /UG=Hs.34578 alpha2,3-sialyltransferase	AK001922		NP_006091
213357_at	0.004341	Homo sapiens, Similar to CG14037 gene product, clone IMAGE:3640720, mRNA, partial cds	AV701318	Hs.356224	
213360_s_at	0.007779	Williams Beuren syndrome chromosome region 20C	AA514622	Hs.295112	NP_742017
213361_at	0.00161	tudor repeat associator with PCTAIRE 2	AW129593	Hs.283761	NP_055105
213365_at	0.033491	KIAA1504 protein	N64622	Hs.157426	
213366_x_at	0.041055	ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	AV711183	Hs.155433	NP_005165
213374_x_at	0.041055	3-hydroxyisobutyryl-Coenzyme A hydrolase	AW000964	Hs.236642	NP_055177
213379_at	1.26E-04	Consensus includes gb:AF091086.1 /DEF=Homo sapiens clone 640 unknown mRNA, complete sequence. /FEA=mRNA /PROD=unknown /DB_XREF=gi:3860009 /UG=Hs.44563 hypothetical protein /FL=gb:NM_015697.1	AF091086		NP_056512

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213397_x_at	0.035171	ribonuclease, RNase A family, 4	AI761728	Hs.283749	NP_002928
213404_s_at	0.007779	601453992F1 NIH_MGC_66 Homo sapiens cDNA clone IMAGE:3857711 5', mRNA sequence.	BF033683		
213414_s_at	7.58E-04	ribosomal protein S19	BE259729	Hs.298262	NP_001013
213436_at	0.016275	G protein-coupled receptor; Human CB1 cannabinoid receptor (CNR1) gene, complete cds.	U73304		
213440_at	1.58E-04	RAB1A, member RAS oncogene family	AL530264	Hs.79507	NP_004152
213448_at	0.009716	metaxin 1	AI693193	Hs.247551	NP_002446
213453_x_at	0.041055	glyceraldehyde-3-phosphate dehydrogenase	BF689355	Hs.169476	NP_002037
213455_at	0.00161	hypothetical protein BC001096	W87466	Hs.352406	NP_612398
213463_s_at	0.041795	Consensus includes gb:AW300504 /FEA=EST /DB_XREF=gi:6710181 /DB_XREF=est:xs66c11.x1 /CLONE=IMAGE:2774612 /UG=Hs.44131 KIAA0974 protein	AB023191		
213465_s_at	5.03E-04	protein phosphatase 1, regulatory subunit 7	BF718769	Hs.36587	NP_002703
213478_at	8.84E-04	Consensus includes gb:AB028949.1 /DEF=Homo sapiens mRNA for KIAA1026 protein, partial cds. /FEA=mRNA /GEN=KIAA1026 /PROD=KIAA1026 protein /DB_XREF=gi:5689388 /UG=Hs.27742 KIAA1026 protein	AB028949		
213501_at	3.26E-04	acyl-Coenzyme A oxidase 1, palmitoyl	T62985	Hs.379991	NP_009223
213508_at	0.016282	ESTs, Weakly similar to R3HU6 ribosomal protein S6, cytosolic - human [H.sapiens]	AA142942	Hs.356665	
213510_x_at	7.58E-04	TL132 protein	AW194543	Hs.234573	NP_665808
213511_s_at	0.016934	Homo sapiens, clone MGC:17492 IMAGE:3453013, mRNA, complete cds	AI167164	Hs.347187	NP_789746
213513_x_at	0.00161	actin related protein 2/3 complex, subunit 2, 34kDa	BG034239	Hs.83583	NP_690601
213515_x_at	1.41E-04	hemoglobin, gamma A	AI133353	Hs.283108	NP_000175
213517_at	6.68E-04	poly(rC) binding protein 2	AW103422	Hs.63525	NP_114366
213521_at	0.02145	ESTs	AW575379	Hs.356456	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213524_s_at	0.007258	Consensus includes gb:NM_015714.1 /DEF=Homo sapiens putative lymphocyte G0G1 switch gene (G0S2), mRNA. /FEA=CDS /GEN=G0S2 /PROD=putative lymphocyte G0G1 switch gene /DB_XREF=gi:7657103 /UG=Hs.95910 putative lymphocyte G0G1 switch gene /FL=gb:NM_015714.1	NM_015714		NP_056529
213526_s_at	0.013223	uncharacterized hematopoietic stem/progenitor cells protein MDS033	BF215644	Hs.54960	NP_061977
213532_at	3.26E-04	a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	AI797833	Hs.380136	
213535_s_at	0.013223	ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast)	AA910614	Hs.84285	NP_003336
213539_at	0.02145	Consensus includes gb:NM_000732.1 /DEF=Homo sapiens CD3D antigen, delta polypeptide (TiT3 complex) (CD3D), mRNA. /FEA=CDS /GEN=CD3D /PROD=CD3D antigen, delta polypeptide (TiT3 complex) /DB_XREF=gi:4502668 /UG=Hs.95327 CD3D antigen, delta polypeptide (TiT3 complex) /FL=gb:NM_000732.1	NM_000732		NP_000723
213541_s_at	0.004659	ESTs	AI351043	Hs.391794	NP_004440
213545_x_at	1.26E-04	sorting nexin 3	BE962615	Hs.12102	NP_690041
213549_at	1.26E-04	ESTs, Moderately similar to PRO0478 protein [Homo sapiens] [H.sapiens]	AI890972	Hs.355844	
213554_s_at	0.010205	hypothetical protein H41	BG257762	Hs.283690	NP_060018
213557_at	0.00228	ESTs	AW305119	Hs.374965	
213567_at	0.003171	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	BF431965	Hs.409065	
213571_s_at	0.00161	eukaryotic translation initiation factor 4E-like 3	BF516289	Hs.19122	NP_004837
213572_s_at	1.26E-04	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1	AI554300	Hs.183583	NP_109591
213574_s_at	2.05E-04	karyopherin (importin) beta 1	AA861608	Hs.180446	NP_002256
213578_at	0.019761	bone morphogenetic protein receptor, type IA	AI678679	Hs.2534	NP_004320
213579_s_at	0.016934	E1A binding protein p300	AI459462	Hs.25272	NP_001420

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213582_at	0.021448	ATPase, Class VI, type 11A	BF439472	Hs.29189	
213588_x_at	0.033381	ribosomal protein L14	AA838274	Hs.738	NP_003964
213593_s_at	6.68E-04	transformer-2 alpha (htra-2 alpha)	AW978896	Hs.24937	
213603_s_at	0.007779	ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2)	BE138888	Hs.301175	NP_002863
213604_at	0.013223	ESTs, Highly similar to FINC_HUMAN Fibronectin precursor (FN) (Cold-insoluble globulin) (CIG) [H.sapiens]	AW451236	Hs.409286	
213614_x_at	0.016934	eukaryotic translation elongation factor 1 alpha 1	BE786672	Hs.181165	
213617_s_at	0.021312	DKFZP586M1523 protein	BF063896	Hs.22981	NP_056291
213619_at	0.004341	heterogeneous nuclear ribonucleoprotein H1 (H)	AV753392	Hs.245710	NP_005511
213620_s_at	0.011626	intercellular adhesion molecule 2	AA126728	Hs.347326	
213623_at	0.004341	Consensus includes gb:NM_007054.1 /DEF=Homo sapiens kinesin family member 3A (KIF3A), mRNA. /FEA=CDS /GEN=KIF3A /PROD=kinesin family member 3A /DB_XREF=gi:6857803 /UG=Hs.43670 kinesin family member 3A /FL=gb:AF041853.1 gb:NM_007054.1	NM_007054		NP_008985
213624_at	3.26E-04	acid sphingomyelinase-like phosphodiesterase	AA873600	Hs.391644	NP_006705
213632_at	0.01436	Consensus includes gb:M94065.1 /DEF=Human dihydroorotate dehydrogenase mRNA, 3 end. /FEA=mRNA /PROD=dihydroorotate dehydrogenase /DB_XREF=gi:555593 /UG=Hs.94925 dihydroorotate dehydrogenase	M94065		NP_001352
213637_at	0.005852	ATP-dependent RNA helicase	BE503392	Hs.380430	NP_689513
213642_at	1.26E-04	ribosomal protein L27	BE312027	Hs.111611	NP_000979
213646_x_at	7.58E-04	tubulin, alpha, ubiquitous	BE300252	Hs.334842	NP_006073
213649_at	2.05E-04	splicing factor, arginine/serine-rich 7, 35kDa	AA524053	Hs.184167	NP_006267
213666_at	6.68E-04	Consensus includes gb:AK026589.1 /DEF=Homo sapiens cDNA: FLJ22936 fis, clone KAT07936. /FEA=mRNA /DB_XREF=gi:10439473 /UG=Hs.90998 KIAA0128 protein; septin 2	AK026589		NP_665801
213671_s_at	0.005852	methionine-tRNA synthetase	AA621558	Hs.279946	NP_004981

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213672_at	0.004541	methionine-tRNA synthetase	AA621558	Hs.279946	NP_004981
213687_s_at	0.007779	ribosomal protein L35a	BE968801	Hs.287361	NP_000987
213694_at	0.016934	ESTs	AW027347	Hs.394035	
213698_at	0.013223	zinc finger protein 258	AI805560	Hs.301637	NP_660353
213702_x_at	1.26E-04	N-acylsphingosine amidohydrolase (acid ceramidase) 1	AI934569	Hs.75811	NP_808592
213708_s_at	7.58E-04	transcription factor-like 4	N40555	Hs.78185	NP_057640
213720_s_at	0.021435	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AI831675	Hs.78202	NP_003063
213726_x_at	3.26E-04	tubulin, beta, 2	AA515698	Hs.251653	
213729_at	2.05E-04	formin binding protein 3	Z78308	Hs.107213	NP_060362
213735_s_at	2.05E-04	cytochrome c oxidase subunit Vb	AI557312	Hs.1342	NP_001853
213741_s_at	7.58E-04	karyopherin alpha 1 (importin alpha 5)	BF575685	Hs.343581	NP_002255
213786_at	0.026891	nudix (nucleoside diphosphate linked moiety X)-type motif 1	AI935415	Hs.153357	NP_006015
213794_s_at	0.041055	DKFZP564O092 protein	AI269117	Hs.9043	
213798_s_at	0.007779	adenylyl cyclase-associated protein	AA806142	Hs.104125	NP_006358
213812_s_at	0.02145	Consensus includes gb:AK024748.1 /DEF=Homo sapiens cDNA: FLJ21095 fis, clone CAS03846. /FEA=mRNA /DB_XREF=gi:10437113 /UG=Hs.108708 calciumcalmodulin-dependent protein kinase kinase 2, beta	AK024748		NP_757380
213817_at	1.26E-04	Consensus includes gb:AL049435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B0220 (from clone DKFZp586B0220). /FEA=mRNA /DB_XREF=gi:4500218 /UG=Hs.170056 Homo sapiens mRNA; cDNA DKFZp586B0220 (from clone DKFZp586B0220)	AL049435		
213826_s_at	0.033381	H3 histone, family 3A	AA292281	Hs.181307	NP_002098

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213831_at	0.022752	Consensus includes gb:X00452.1 /DEF=Human mRNA for DC classII histocompatibility antigen alpha-chain. /FEA=mRNA /PROD=DC classII histocompatibility antigenalpha-chain /DB_XREF=gi:32265 /UG=Hs.198253 major histocompatibility complex, class II, DQ alpha 1	X00452		NP_002113
213836_s_at	1.26E-04	KIAA1001 protein	AW052084	Hs.279937	NP_055775
213850_s_at	6.68E-04	splicing factor, arginine/serine-rich 2, interacting protein	AI984932	Hs.51957	NP_004710
213851_at	0.001012	ESTs, Weakly similar to S65173 probable membrane protein YPL162c - yeast (Saccharomyces cerevisiae) [S.cerevisiae]	BG031677	Hs.394011	
213860_x_at	2.05E-04	casein kinase 1, alpha 1	AW268585	Hs.283738	NP_001883
213867_x_at	0.013223	actin, beta	AA809056	Hs.288061	NP_001092
213882_at	0.003529	beta-amyloid binding protein precursor	AA012917	Hs.333541	NP_114416
213883_s_at	0.003171	beta-amyloid binding protein precursor	AA012917	Hs.333541	NP_114416
213902_at	0.00228	N-acylsphingosine amidohydrolase (acid ceramidase) 1	AI379338	Hs.75811	NP_808592
213911_s_at	2.05E-04	H2A histone family, member Z	BF718636	Hs.119192	NP_002097
213918_s_at	5.03E-04	IDN3 protein	BF221673	Hs.225767	NP_597677
213919_at	0.034221	DnaJ (Hsp40) homolog, subfamily C, member 4	AW024467	Hs.172847	NP_005519
213922_at	0.003539	KIAA0847 protein	AW294686	Hs.356431	NP_775771
213923_at	5.03E-04	RAP2B, member of RAS oncogene family	AW005535	Hs.388235	NP_002877
213936_x_at	0.022795	ESTs, Moderately similar to hypothetical protein FLJ20294 [Homo sapiens] [H.sapiens]	AW276646	Hs.355462	
213940_s_at	0.003171	formin-binding protein 17	AU145053	Hs.408992	
213956_at	6.68E-04	centrosome-associated protein 350	AW299294	Hs.92200	NP_055625
213958_at	0.001618	CD6 antigen	AW134823	Hs.81226	NP_006716
213969_x_at	0.004341	ribosomal protein L29	BF683426	Hs.350068	
213971_s_at	0.022708	joined to JAZF1	AI924660	Hs.197803	NP_056170
213988_s_at	3.26E-04	spermidine/spermine N1-acetyltransferase	BE971383	Hs.396709	NP_002961
213995_at	0.02145	mitochondrial ATP synthase regulatory component factor B	AW195882	Hs.56155	NP_056499
214001_x_at	0.004341	ribosomal protein S10	AA320764	Hs.76230	NP_001005

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214022_s_at	0.041055	interferon induced transmembrane protein 1 (9-27)	AA749101	Hs.146360	
214032_at	6.68E-04	ESTs, Highly similar to ZA70_HUMAN Tyrosine-protein kinase ZAP-70 (70 kDa zeta-associated protein) (Syk-related tyrosine kinase) [H.sapiens]	AI817942	Hs.406272	
214033_at	0.008237	ATP-binding cassette, sub-family C (CFTR/MRP), member 6	AI084637	Hs.274260	NP_001162
214034_at	0.014208	type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	BE551138	Hs.247043	NP_057526
214039_s_at	0.027633	putative integral membrane transporter	T15777	Hs.296398	NP_060877
214041_x_at	2.05E-04	ribosomal protein L37a	BE857772	Hs.296290	NP_000989
214055_x_at	6.68E-04	KIAA1096 protein	AW238632	Hs.69559	NP_055987
214080_x_at	0.004341	protein kinase C substrate 80K-H	AI815793	Hs.1432	NP_002734
214081_at	0.040064	Consensus includes gb:AF070526.1 /DEF=Homo sapiens clone 24787 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387881 /UG=Hs.13429 Homo sapiens clone 24787 mRNA sequence	AF070526		NP_065138
214085_x_at	0.02145	glioma pathogenesis-related protein	AI912583	Hs.154762	NP_008974
214088_s_at	0.004452	fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group included)	AW080549	Hs.169238	NP_000140
214112_s_at	1.26E-04	Homo sapiens, clone MGC:10039 IMAGE:3889701, mRNA, complete cds	AA543076	Hs.82171	NP_835225
214119_s_at	0.033381	FK506 binding protein 1A, 12kDa	AI936769	Hs.380080	NP_463460
214121_x_at	0.007942	enigma (LIM domain protein)	AA086229	Hs.102948	NP_005442
214136_at	0.011788	nudix (nucleoside diphosphate linked moiety X)-type motif 13	W80642	Hs.16951	
214151_s_at	1.26E-04	phosphatidylinositol glycan, class B	AU144243	Hs.247118	NP_004846
214152_at	3.26E-04	phosphatidylinositol glycan, class B	AU144243	Hs.247118	NP_004846
214153_at	0.005411	homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2	BE467941	Hs.408991	
214173_x_at	0.03018	chromosome 19 open reading frame 2	AW514900	Hs.7943	NP_604431
214177_s_at	0.010205	hematopoietic PBX-interacting protein	AI935162	Hs.8068	NP_065385
214181_x_at	0.033381	leukocyte specific transcript 1	AI735692	Hs.88411	NP_009092
214198_s_at	3.26E-04	DiGeorge syndrome critical region gene 2	AU150824	Hs.2491	NP_005128

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214211_at	0.041055	ferritin, heavy polypeptide 1	AA083483	Hs.62954	NP_004174
214224_s_at	0.041055	protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE674061	Hs.11774	NP_006214
214246_x_at	0.016934	Misshapen/NIK-related kinase	AI859060	Hs.112028	NP_000071
214257_s_at	0.016934	SEC22 vesicle trafficking protein-like 1 (<i>S. cerevisiae</i>)	AA890010	Hs.50785	NP_004883
214259_s_at	0.026891	aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)	AW074911	Hs.6980	NP_003680
214266_s_at	0.046329	enigma (LIM domain protein)	AW206786	Hs.102948	NP_005442
214268_s_at	0.02145	myotubularin related protein 4	AL042220	Hs.141727	NP_004678
214273_x_at	0.039149	Conserved gene telomeric to alpha globin cluster	AV704353	Hs.19699	
214274_s_at	0.007779	acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase)	AI860341	Hs.166160	NP_001598
214280_x_at	0.013223	Consensus includes gb:X79536.1 /DEF=H.sapiens mRNA for hnRNPcore protein A1. /FEA=mRNA /PROD=hnRNPcore protein A1 /DB_XREF=gi:496897 /UG=Hs.249495 heterogeneous nuclear ribonucleoprotein A1	X79536		NP_112420
214281_s_at	7.58E-04	zinc finger protein 363	AA524525	Hs.48297	NP_056251
214288_s_at	0.00161	proteasome (prosome, macropain) subunit, beta type, 1	W86293	Hs.75748	NP_002784
214290_s_at	0.041055	H2A histone family, member O	AA451996	Hs.795	NP_003507
214299_at	0.021435	topoisomerase (DNA) III alpha	AI676092	Hs.91175	NP_004609
214305_s_at	6.68E-04	splicing factor 3b, subunit 1, 155kDa	AW003030	Hs.334826	NP_036565
214315_x_at	0.005852	calreticulin	AI348935	Hs.16488	NP_004334
214329_x_at	0.013223	tumor necrosis factor (ligand) superfamily, member 10	AW474434	Hs.83429	NP_003801
214352_s_at	3.26E-04	v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	BF673699	Hs.351221	NP_203524
214366_s_at	6.68E-04	arachidonate 5-lipoxygenase	AA995910	Hs.89499	NP_000689
214403_x_at	1.72E-04	prostate epithelium-specific Ets transcription factor	AI307915	Hs.79414	NP_036523
214405_at	0.018693	ESTs	Z39557	Hs.389152	
214414_x_at	0.026891	hemoglobin, alpha 2	T50399	Hs.347939	NP_000549
214429_at	2.05E-04	Consensus includes gb:U47635.1 /DEF=Human D13S824E locus mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:1669390 /UG=Hs.79877 myotubularin related protein 6 /FL=gb:U47635.1	U47635		NP_004676

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214430_at	7.58E-04	Consensus includes gb:NM_000169.1 /DEF=Homo sapiens galactosidase, alpha (GLA), mRNA. /FEA=CDS /GEN=GLA /PROD=galactosidase, alpha /DB_XREF=gi:4504008 /UG=Hs.69089 galactosidase, alpha /FL=gb:BC002689.1 gb:NM_000169.1	NM_000169		NP_000160
214438_at	0.041055	Consensus includes gb:M60721.1 /DEF=Human homeobox gene, complete cds. /FEA=mRNA /DB_XREF=gi:183789 /UG=Hs.74870 H2.0 (Drosophila)-like homeo box 1 /FL=gb:NM_021958.1 gb:M60721.1	M60721		NP_068777
214439_x_at	0.0174	Consensus includes gb:AF043899.1 /DEF=Homo sapiens amphiphysin IIc1 mRNA, complete cds. /FEA=CDS /PROD=amphiphysin IIc1 /DB_XREF=gi:3064256 /UG=Hs.193163 bridging integrator 1 /FL=gb:AF043899.1	AF043899		NP_647601
214440_at	7.58E-04	Consensus includes gb:NM_000662.1 /DEF=Homo sapiens N-acetyltransferase 1 (arylamine N-acetyltransferase) (NAT1), mRNA. /FEA=CDS /GEN=NAT1 /PROD=N-acetyltransferase 1 /DB_XREF=gi:4505334 /UG=Hs.155956 N-acetyltransferase 1 (arylamine N-acetyltransferase) /FL=gb:NM_000662.1	NM_000662		NP_000653

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214470_at	0.041055	Consensus includes gb:NM_002258.1 /DEF=Homo sapiens killer cell lectin-like receptor subfamily B, member 1 (KLRB1), mRNA. /FEA=CDS /GEN=KLRB1 /PROD=killer cell lectin-like receptor subfamily B,member 1 /DB_XREF=gi:4504878 /UG=Hs.169824 killer cell lectin-like receptor subfamily B, member 1 /FL=gb:NM_002258.1 gb:U11276.1	NM_002258		NP_002249
214473_x_at	0.033381	Consensus includes gb:NM_005395.1 /DEF=Homo sapiens postmeiotic segregation increased 2-like 9 (PMS2L9), mRNA. /FEA=CDS /GEN=PMS2L9 /PROD=postmeiotic segregation increased 2-like 9 /DB_XREF=gi:4885552 /UG=Hs.301862 postmeiotic segregation increased 2-like 9 /FL=gb:NM_005395.1	NM_005395		NP_005386
214496_x_at	5.64E-04	Consensus includes gb:NM_012330.1 /DEF=Homo sapiens histone acetyltransferase (MORF), mRNA. /FEA=CDS /GEN=MORF /PROD=histone acetyltransferase /DB_XREF=gi:6912511 /UG=Hs.27590 histone acetyltransferase /FL=gb:AF119231.1 gb:NM_012330.1	NM_012330		NP_036462
214501_s_at	5.03E-04	Consensus includes gb:AF044286.1 /DEF=Homo sapiens histone macroH2A1.1 mRNA, complete cds. /FEA=CDS /PROD=histone macroH2A1.1 /DB_XREF=gi:3493530 /UG=Hs.75258 H2A histone family, member Y /FL=gb:AF044286.1	AF044286		NP_613258

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214511_x_at	7.58E-04	Consensus includes gb:L03419.1 /DEF=Human Fc-gamma receptor I B1 mRNA, complete cds. /FEA=CDS /PROD=Fc gamma receptor I /DB_XREF=gi:182460 /UG=Hs.77424 Fc fragment of IgG, high affinity Ia, receptor for (CD64) /FL=gb:L03419.1 gb:L03420.1	L03419		NP_000557
214512_s_at	0.00228	Consensus includes gb:NM_006713.1 /DEF=Homo sapiens activated RNA polymerase II transcription cofactor 4 (PC4), mRNA. /FEA=CDS /GEN=PC4 /PROD=activated RNA polymerase II transcriptioncofactor 4 /DB_XREF=gi:5729967 /UG=Hs.74861 activated RNA polymerase II transcription cofactor 4 /FL=gb:U12979.1 gb:NM_006713.1	NM_006713		NP_006704
214523_at	3.93E-04	Consensus includes gb:NM_001805.1 /DEF=Homo sapiens CCAATenhancer binding protein (CEBP), epsilon (CEBPE), mRNA. /FEA=CDS /GEN=CEBPE /PROD=CCAATenhancer binding protein (CEBP), epsilon /DB_XREF=gi:4502766 /UG=Hs.158323 CCAATenhancer binding protein (CEBP), epsilon /FL=gb:NM_001805.1	NM_001805		NP_001796
214531_s_at	2.05E-04	Consensus includes gb:AF065484.1 /DEF=Homo sapiens sorting nexin 1A (SNX1) mRNA, complete cds. /FEA=CDS /GEN=SNX1 /PROD=sorting nexin 1A /DB_XREF=gi:3152941 /UG=Hs.75283 sorting nexin 1 /FL=gb:AF065484.1	AF065484		NP_690039

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214548_x_at	1.26E-04	Consensus includes gb:AF064092.1 /DEF=Homo sapiens mutant guanine nucleotide-binding protein G(s), alpha subunit mRNA, complete cds. /FEA=CDS /PROD=mutant guanine nucleotide-binding protein G(s),alpha subunit /DB_XREF=gi:3142636 /UG=Hs.273385 guanine nucleotide binding protein (G protein), alpha stimulating activity polypeptide 1 /FL=gb:AF064092.1	AF064092		NP_536351
214553_s_at	0.003862	Consensus includes gb:NM_006628.1 /DEF=Homo sapiens cyclic AMP phosphoprotein, 19 kD (ARPP-19), mRNA. /FEA=CDS /GEN=ARPP-19 /PROD=cyclic AMP phosphoprotein, 19 kD /DB_XREF=gi:5729731 /UG=Hs.7351 cyclic AMP phosphoprotein, 19 kD /FL=gb:NM_006628.1	NM_006628		NP_006619
214561_at	0.027243	Consensus includes gb:NM_024317.1 /DEF=Homo sapiens immunoglobulin-like transcript 10 (ILT10), mRNA. /FEA=CDS /GEN=ILT10 /PROD=immunoglobulin-like transcript 10 /DB_XREF=gi:13399319 /UG=Hs.202680 immunoglobulin-like transcript 10 /FL=gb:NM_024317.1	NM_024317		NP_077293
214572_s_at	0.038947	Consensus includes gb:NM_005543.1 /DEF=Homo sapiens insulin-like 3 (Leydig cell) (INSL3), mRNA. /FEA=CDS /GEN=INSL3 /PROD=insulin-like 3 (Leydig cell) /DB_XREF=gi:5031802 /UG=Hs.37062 insulin-like 3 (Leydig cell) /FL=gb:NM_005543.1	NM_005543		NP_005534

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214574_x_at	0.007779	Consensus includes gb:NM_007161.1 /DEF=Homo sapiens DNA segment on chromosome 6 (unique) 49 expressed sequence, NK cell triggering receptor, p30 (D6S49E), mRNA. /FEA=CDS /GEN=D6S49E /PROD=leukocyte-specific transcript 1 /DB_XREF=gi:6005740 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:NM_007161.1	NM_007161		NP_009092
214583_at	0.012367	regulatory solute carrier protein, family 1, member 1	AI268381	Hs.239459	NP_006502
214590_s_at	2.05E-04	ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)	AL545760	Hs.129683	NP_003329
214594_x_at	0.005852	ATPase, Class I, type 8B, member 1	BG252666	Hs.388624	NP_005594
214606_at	0.001303	tetraspan 2	BF129969	Hs.234863	NP_005716
214615_at	0.003539	Consensus includes gb:NM_014499.1 /DEF=Homo sapiens putative purinergic receptor (P2Y10), mRNA. /FEA=CDS /GEN=P2Y10 /PROD=putative purinergic receptor /DB_XREF=gi:10092632 /UG=Hs.296433 putative purinergic receptor /FL=gb:NM_014499.1	NM_014499		NP_055314
214617_at	0.004341	perforin 1 (pore forming protein)	AI445650	Hs.2200	NP_005032
214620_x_at	3.56E-04	peptidylglycine alpha-amidating monooxygenase	BF038548	Hs.83920	NP_620177
214629_x_at	0.001116	Consensus includes gb:AF320999.1 /DEF=Homo sapiens Nogo-A protein short form mRNA, complete cds, alternatively spliced. /FEA=CDS /PROD=Nogo-A protein short form /DB_XREF=gi:11878277 /UG=Hs.65450 reticulon 4 /FL=gb:AF320999.1	AF320999		NP_722550

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214651_s_at	0.011725	Consensus includes gb:U41813.1 /DEF=Human class I homeoprotein (HOXA9) mRNA, partial cds. /FEA=mRNA /GEN=HOXA9 /PROD=HOXA9 /DB_XREF=gi:1184168 /UG=Hs.127428 homeo box A9 /FL=gb:NM_002142.1	U41813		NP_689952
214658_at	0.00228	CGI-109 protein	BG286537	Hs.278391	
214663_at	0.00228	Consensus includes gb:AB007941.1 /DEF=Homo sapiens mRNA for KIAA0472 protein, partial cds. /FEA=mRNA /GEN=KIAA0472 /PROD=KIAA0472 protein /DB_XREF=gi:3413905 /UG=Hs.6874 KIAA0472 protein	AB007941		
214665_s_at	1.26E-04	Consensus includes gb:AK000095.1 /DEF=Homo sapiens cDNA FLJ20088 fis, clone COL03869. /FEA=mRNA /DB_XREF=gi:7019960 /UG=Hs.85301 calcium binding protein P22	AK000095		
214669_x_at	0.007052	immunoglobulin kappa constant	BG485135	Hs.306357	
214684_at	5.03E-04	Consensus includes gb:X63381.1 /DEF=Homo sapiens mRNA for serum response factor-related protein, RSRFC4. /FEA=mRNA /GEN=RSRFC4 /PROD=serum response factor-related protein /DB_XREF=gi:432655 /UG=Hs.182280 MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	X63381		NP_005578
214687_x_at	0.033381	Consensus includes gb:AK026577.1 /DEF=Homo sapiens cDNA: FLJ22924 fis, clone KAT06977, highly similar to HSALDAR Human fibroblast mRNA for aldolase A. /FEA=mRNA /DB_XREF=gi:10439461 /UG=Hs.273415 aldolase A, fructose-bisphosphate	AK026577		NP_000025
214691_x_at	0.002704	hypothetical protein KIAA1164	AU121431	Hs.40719	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214693_x_at	0.029453	hypothetical protein DJ328E19.C1.1	BE732345	Hs.323463	NP_056198
214696_at	0.033381	Consensus includes gb:AF070569.1 /DEF=Homo sapiens clone 24659 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387938 /UG=Hs.29206 Homo sapiens clone 24659 mRNA sequence	AF070569		NP_116284
214703_s_at	7.58E-04	KIAA0935 protein	AW954107	Hs.12183	
214714_at	0.003741	Consensus includes gb:AK022360.1 /DEF=Homo sapiens cDNA FLJ12298 fis, clone MAMMA1001837, weakly similar to ZINC FINGER PROTEIN 29. /FEA=mRNA /DB_XREF=gi:10433741 /UG=Hs.284168 hypothetical protein MPMGp800M05499Q3	AK022360		NP_115540
214715_x_at	0.016934	Consensus includes gb:AK024789.1 /DEF=Homo sapiens cDNA: FLJ21136 fis, clone CAS07469. /FEA=mRNA /DB_XREF=gi:10437175 /UG=Hs.206882 Homo sapiens mRNA for FLJ00032 protein, partial cds	AK024789		NP_150630
214718_at	6.68E-04	Consensus includes gb:AK026142.1 /DEF=Homo sapiens cDNA: FLJ22489 fis, clone HRC10951. /FEA=mRNA /DB_XREF=gi:10438897 /UG=Hs.21145 hypothetical protein FLJ22489	AK026142		NP_066990
214719_at	0.010205	Consensus includes gb:AK026720.1 /DEF=Homo sapiens cDNA: FLJ23067 fis, clone LNG04993. /FEA=mRNA /DB_XREF=gi:10439638 /UG=Hs.117167 Homo sapiens cDNA: FLJ23067 fis, clone LNG04993	AK026720		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214730_s_at	0.013223	Consensus includes gb:AK025457.1 /DEF=Homo sapiens cDNA: FLJ21804 fis, clone HEP00746, highly similar to HSU64791 Human Golgi membrane sialoglycoprotein MG160 (GLG1) mRNA. /FEA=mRNA /DB_XREF=gi:10437977 /UG=Hs.78979 Golgi apparatus protein 1	AK025457		NP_036333
214733_s_at	4.42E-04	Consensus includes gb:AL031427 /DEF=Human DNA sequence from clone 167A19 on chromosome 1p32.1-33. Contains three genes for novel proteins, the DIO1 gene for type I iodothyronine deiodinase (EC 3.8.1.4, TXDI1, ITDI1) and an HNRNP A3 (Heterogenous Nuclear Ribonucleoprotein A3, FBRNP) ... /FEA=mRNA_6 /DB_XREF=gi:4835258 /UG=Hs.11923 hypothetical protein	AL031427		
214737_x_at	0.005852	heterogeneous nuclear ribonucleoprotein C (C1/C2)	AV725195	Hs.182447	NP_112604
214743_at	0.003171	ESTs, Highly similar to CUT1_HUMAN CCAAT displacement protein (CDP) (Cut-like 1) [H.sapiens]	BE046521	Hs.381906	NP_001904
214748_at	0.016934	Consensus includes gb:U50529.1 /DEF=Human BRCA2 region, mRNA sequence CG016. /FEA=mRNA /DB_XREF=gi:1531600 /UG=Hs.112434 Novel human gene mapping to chromosome 13	U50529		NP_149102
214766_s_at	0.013223	Consensus includes gb:AL080144.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N093 (from clone DKFZp434N093); partial cds. /FEA=mRNA /GEN=DKFZp434N093 /PROD=hypothetical protein /DB_XREF=gi:5262592 /UG=Hs.33363 DKFZP434N093 protein	AL080144		NP_056261

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214773_x_at	0.009689	similar to CG9578 gene product	AI983505	Hs.137576	NP_690866
214783_s_at	0.005852	annexin A11	BG177920	Hs.75510	NP_665876
214785_at	0.005852	Consensus includes gb:AB023203.1 /DEF=Homo sapiens mRNA for KIAA0986 protein, partial cds. /FEA=mRNA /GEN=KIAA0986 /PROD=KIAA0986 protein /DB_XREF=gi:4589615 /UG=Hs.53542 KIAA0986 protein	AB023203		NP_150648
214791_at	0.033381	Consensus includes gb:AK023116.1 /DEF=Homo sapiens cDNA FLJ13054 fis, clone NT2RP3001527, highly similar to Human Sp140 protein (Sp140) mRNA. /FEA=mRNA /DB_XREF=gi:10434889 /UG=Hs.158761 Homo sapiens cDNA FLJ13054 fis, clone NT2RP3001527, highly similar to Human Sp140 protein (Sp140) mRNA	AK023116		NP_612411
214798_at	0.006698	KIAA0703 gene product	AW291664	Hs.6168	
214801_at	0.004798	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	W88821	Hs.408958	
214805_at	2.05E-04	Consensus includes gb:U79273.1 /DEF=Human clone 23933 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710239 /UG=Hs.239483 Human clone 23933 mRNA sequence	U79273		
214808_at	0.001747	ESTs, Moderately similar to 2211404A B219/OB receptor [Homo sapiens] [H.sapiens]	AU147851	Hs.194397	
214812_s_at	3.26E-04	Consensus includes gb:D80006.1 /DEF=Human mRNA for KIAA0184 gene, partial cds. /FEA=mRNA /GEN=KIAA0184 /DB_XREF=gi:1136427 /UG=Hs.322903 KIAA0184 protein	D80006		
214813_at	0.001012	ESTs, Moderately similar to A48913 KRAB zinc finger protein 75 - human [H.sapiens]	W90796	Hs.389646	NP_009062
214814_at	0.007052	KIAA1966 protein	BF592058	Hs.158184	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214816_x_at	0.021244	Consensus includes gb:BC003535.1 /DEF=Homo sapiens, clone IMAGE:3609326, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3609326) /DB_XREF=gi:13097635 /UG=Hs.192127 Homo sapiens, clone IMAGE:3609326, mRNA, partial cds	BC003535		NP_689479
214843_s_at	0.003529	Consensus includes gb:AK022864.1 /DEF=Homo sapiens cDNA FLJ12802 fis, clone NT2RP2002124, weakly similar to UBIQUITIN CARBOXYL TERMINAL HYDROLASE 4 (EC 3.1.2.15). /FEA=mRNA /DB_XREF=gi:10434503 /UG=Hs.173694 KIAA1097 protein	AK022864		NP_055832
214848_at	0.036254	Consensus includes gb:U79277.1 /DEF=Human clone 23548 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710245 /UG=Hs.71848 Human clone 23548 mRNA sequence	U79277		
214853_s_at	0.003171	SHC (Src homology 2 domain containing) transforming protein 1	AI091079	Hs.81972	NP_003020
214857_at	0.00884	Consensus includes gb:AL050035.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566H0124 (from clone DKFZp566H0124). /FEA=mRNA /DB_XREF=gi:4884276 /UG=Hs.133130 Homo sapiens mRNA; cDNA DKFZp566H0124 (from clone DKFZp566H0124)	AL050035		
214866_at	0.001911	Consensus includes gb:X74039.1 /DEF=H.sapiens mRNA for urokinase plasminogen activator receptor. /FEA=mRNA /PROD=urokinase plasminogen activator receptor /DB_XREF=gi:456192 /UG=Hs.179657 plasminogen activator, urokinase receptor	X74039		NP_002650
214875_x_at	0.016934	amyloid beta (A4) precursor-like protein 2	AW001847	Hs.279518	NP_001633

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214895_s_at	0.010205	a disintegrin and metalloproteinase domain 10	AU135154	Hs.172028	NP_001101
214902_x_at	0.007779	Consensus includes gb:AL080232.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A061 (from clone DKFZp586A061). /FEA=mRNA /DB_XREF=gi:5262725 /UG=Hs.220696 Homo sapiens mRNA; cDNA DKFZp586A061 (from clone DKFZp586A061)	AL080232		
214906_x_at	0.004341	Consensus includes gb:AL049786.1 /DEF=Novel human gene mapping to chromosome 13. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:4902708 /UG=Hs.22174 Novel human gene mapping to chromosome 13	AL049786		NP_438169
214909_s_at	0.005339	Consensus includes gb:AK026191.1 /DEF=Homo sapiens cDNA: FLJ22538 fis, clone HRC13212, highly similar to AF070667 Homo sapiens NG,NG-dimethylarginine dimethylaminohydrolase homolog mRNA. /FEA=mRNA /DB_XREF=gi:10438961 /UG=Hs.247362 dimethylarginine dimethylaminohydrolase 2	AK026191		NP_039268
214911_s_at	6.68E-04	Consensus includes gb:S78771.1 /DEF=NAT=CpG island-associated gene human, mRNA, 1741 nt. /FEA=mRNA /DB_XREF=gi:244232 /UG=Hs.75243 bromodomain-containing 2	S78771		NP_005095
214915_at	0.005339	Consensus includes gb:AK021842.1 /DEF=Homo sapiens cDNA FLJ11780 fis, clone HEMBA1005931, weakly similar to ZINC FINGER PROTEIN 83. /FEA=mRNA /DB_XREF=gi:10433110 /UG=Hs.279935 Homo sapiens cDNA FLJ11780 fis, clone HEMBA1005931, weakly similar to ZINC FINGER PROTEIN 83	AK021842		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214917_at	6.79E-04	Consensus includes gb:AK024252.1 /DEF=Homo sapiens cDNA FLJ14190 fis, clone NT2RP2006534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC ALPHA-1 CHAIN (EC 2.7.1.-). /FEA=mRNA /DB_XREF=gi:10436581 /UG=Hs.288546 Homo sapiens cDNA FLJ14190 fis, clone NT2RP2006534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC ALPHA-1 CHAIN (EC 2.7.1.-)	AK024252		
214919_s_at	7.58E-04	FLJ20288 protein	R39094	Hs.84045	NP_003723
214938_x_at	2.05E-04	Consensus includes gb:AF283771.2 /DEF=Homo sapiens clone TCBAP0774 mRNA sequence. /FEA=mRNA /DB_XREF=gi:10281740 /UG=Hs.274472 high-mobility group (nonhistone chromosomal) protein 1	AF283771		NP_002119
214947_at	7.10E-04	Consensus includes gb:AF052146.1 /DEF=Homo sapiens clone 24653 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360455 /UG=Hs.155085 Homo sapiens clone 24653 mRNA sequence	AF052146		
214953_s_at	2.05E-04	Consensus includes gb:X06989.1 /DEF=Human mRNA for amyloid A4(751) protein. /FEA=mRNA /DB_XREF=gi:28720 /UG=Hs.177486 amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	X06989		NP_000475
214962_s_at	0.03002	Consensus includes gb:AK026236.1 /DEF=Homo sapiens cDNA: FLJ22583 fis, clone HSI02596. /FEA=mRNA /DB_XREF=gi:10439023 /UG=Hs.22559 KIAA0197 protein	AK026236		
214964_at	0.031948	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AA554430	Hs.269421	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214965_at	0.034221	Consensus includes gb:AF070574.1 /DEF=Homo sapiens clone 24819 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387945 /UG=Hs.22316 Homo sapiens clone 24819 mRNA sequence	AF070574		NP_689552
214972_at	0.0174	meningioma expressed antigen 5 (hyaluronidase)	AU144791	Hs.5734	
214988_s_at	0.016934	Consensus includes gb:X63071.1 /DEF=H.sapiens mRNA for novel DNA binding protein. /FEA=mRNA /GEN=DBP-5 /PROD=DBP-5 protein /DB_XREF=gi:30487 /UG=Hs.92909 SON DNA binding protein	X63071		NP_620305
214992_s_at	0.011725	Consensus includes gb:AD000092 /DEF=Homo sapiens DNA from chromosome 19p13.2 cosmid R31240, R30272 and R28549 containing the EKLF, GCDH, CRTC, and RAD23A genes, genomic sequence /FEA=CDS_1 /DB_XREF=gi:1905905 /UG=Hs.307251 Homo sapiens DNA from chromosome 19p13.2 cosmid R31240, R30272 and R28549 containing the EKLF, GCDH, CRTC, and RAD23A genes, genomic sequence	AD000092		
215000_s_at	0.001116	Consensus includes gb:AL117593.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G082 (from clone DKFZp564G082). /FEA=mRNA /DB_XREF=gi:5912157 /UG=Hs.103419 fasciculation and elongation protein zeta 2 (zygin II)	AL117593		NP_005093
215004_s_at	0.034317	Hypothetical human protein; Similar to novel proteins encoded by (AC004447) KIAA0365 [Homo sapiens] and (Y08265) DAN26 [Homo sapiens]; Homo sapiens chromosome 19, cosmid F23858, complete sequence.	AC004475		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215030_at	0.046651	Consensus includes gb:AK023187.1 /DEF=Homo sapiens cDNA FLJ13125 fis, clone NT2RP3002877. /FEA=mRNA /DB_XREF=gi:10435001 /UG=Hs.287551 Homo sapiens cDNA FLJ13125 fis, clone NT2RP3002877	AK023187		
215031_x_at	0.013088	hypothetical protein FLJ20552	BG420893	Hs.69554	NP_060346
215032_at	0.004341	Consensus includes gb:AK022442.1 /DEF=Homo sapiens cDNA FLJ12380 fis, clone MAMMA1002556. /FEA=mRNA /DB_XREF=gi:10433838 /UG=Hs.300934 Homo sapiens cDNA FLJ12380 fis, clone MAMMA1002556	AK022442		
215038_s_at	7.58E-04	Consensus includes gb:AF049103.1 /DEF=Homo sapiens Huntingtin interacting protein mRNA, partial cds. /FEA=mRNA /PROD=Huntingtin interacting protein /DB_XREF=gi:3319281 /UG=Hs.306184 Huntingtin interacting protein B	AF049103		NP_054878
215043_s_at	0.022988	Consensus includes gb:X83301.1 /DEF=H.sapiens SMA5 mRNA. /FEA=mRNA /GEN=SMA5 /DB_XREF=gi:603029 /UG=Hs.324728 SMA5	X83301		
215047_at	1.58E-04	Consensus includes gb:AL080170.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C091 (from clone DKFZp434C091); partial cds. /FEA=mRNA /GEN=DKFZp434C091 /PROD=hypothetical protein /DB_XREF=gi:5262639 /UG=Hs.51692 DKFZP434C091 protein	AL080170		NP_056246

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215049_x_at	0.005852	Consensus includes gb:Z22969.1 /DEF=H.sapiens mRNA for M130 antigen cytoplasmic variant 1. /FEA=mRNA /PROD=M130 antigen cytoplasmic variant 1 /DB_XREF=gi:312143 /UG=Hs.74076 CD163 antigen	Z22969		NP_004235
215051_x_at	5.03E-04	allograft inflammatory factor 1	BF213829	Hs.76364	NP_116573
215063_x_at	0.001886	Consensus includes gb:AL390149.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547B026 (from clone DKFZp547B026). /FEA=mRNA /DB_XREF=gi:9368883 /UG=Hs.283862 Homo sapiens mRNA; cDNA DKFZp547B026 (from clone DKFZp547B026)	AL390149		
215087_at	7.58E-04	Consensus includes gb:AL109730.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 68600. /FEA=mRNA /DB_XREF=gi:5689835 /UG=Hs.306331 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 68600	AL109730		
215088_s_at	1.26E-04	ESTs, Highly similar to C560_HUMAN Succinate dehydrogenase cytochrome B560 subunit, mitochondrial precursor (QPS1) (CII-3) (Succinate dehydrogenase complex subunit C) (Succinate-ubiquinone oxidoreductase cytochrome B large subunit) (CYBL) [H.sapiens]	BG110532	Hs.355964	
215118_s_at	0.040938	Homo sapiens translocation associated fusion protein IRTA1/IGA1 (IRTA1/IGHA1) mRNA, complete cds	AW519168	Hs.367852	
215127_s_at	2.05E-04	RNA binding motif, single stranded interacting protein 1	AL517946	Hs.241567	NP_058523

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215158_s_at	1.26E-04	Consensus includes gb:AK022531.1 /DEF=Homo sapiens cDNA FLJ12469 fis, clone NT2RM1000874, highly similar to Homo sapiens death effector domain-containing testicular molecule mRNA. /FEA=mRNA /DB_XREF=gi:10433972 /UG=Hs.169681 death effector domain-containing	AK022531		NP_127491
215175_at	3.26E-04	Consensus includes gb:AB023212.1 /DEF=Homo sapiens mRNA for KIAA0995 protein, partial cds. /FEA=mRNA /GEN=KIAA0995 /PROD=KIAA0995 protein /DB_XREF=gi:4589633 /UG=Hs.225967 KIAA0995 protein	AB023212		NP_055797
215176_x_at	0.022708	immunoglobulin kappa constant	AW404894	Hs.306357	
215193_x_at	0.004341	Consensus includes gb:AJ297586.1 /DEF=Homo sapiens mRNA for MHC class II antigen (HLA-DRB1 gene), DRB1*0402 allele. /FEA=CDS /GEN=HLA-DRB1 /PROD=MHC class II antigen /DB_XREF=gi:10185079 /UG=Hs.279930 major histocompatibility complex, class II, DR beta 3	AJ297586		NP_002115
215204_at	0.010205	ESTs, Weakly similar to 2109260A B cell growth factor [Homo sapiens] [H.sapiens]	AU147295	Hs.288575	
215210_s_at	0.003862	Consensus includes gb:S72422.1 /DEF=E2k=alpha-ketoglutarate dehydrogenase complex dihydrolipoyl succinyltransferase human, fetal brain, mRNA, 2987 nt. /FEA=mRNA /GEN=E2k /PROD=alpha-ketoglutarate dehydrogenase complexdihydrolipoyl succinyltransferase /DB_XREF=gi:632883 /UG=Hs.296348 E2k	S72422		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215221_at	1.26E-04	Consensus includes gb:AK025064.1 /DEF=Homo sapiens cDNA: FLJ21411 fis, clone COL03986. /FEA=mRNA /DB_XREF=gi:10437503 /UG=Hs.306758 Homo sapiens cDNA: FLJ21411 fis, clone COL03986	AK025064		
215222_x_at	0.016934	Consensus includes gb:AK023406.1 /DEF=Homo sapiens cDNA FLJ13344 fis, clone OVARC1002050, highly similar to Homo sapiens mRNA for actin binding protein ABP620. /FEA=mRNA /DB_XREF=gi:10435330 /UG=Hs.321395 Homo sapiens cDNA FLJ13344 fis, clone OVARC1002050, highly similar to Homo sapiens mRNA for actin binding protein ABP620	AK023406		NP_149033
215223_s_at	0.00228	superoxide dismutase 2, mitochondrial	W46388	Hs.372783	NP_000627
215236_s_at	0.008185	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)	AV721177	Hs.114765	NP_665801
215273_s_at	0.001054	Consensus includes gb:AK024982.1 /DEF=Homo sapiens cDNA: FLJ21329 fis, clone COL02464. /FEA=mRNA /DB_XREF=gi:10437412 /UG=Hs.158196 transcriptional adaptor 3 (ADA3, yeast homolog) like (PCAF histone acetylase complex)	AK024982		NP_597815
215275_at	0.013223	EST375211 MAGE resequences, MAGH Homo sapiens cDNA, mRNA sequence.	AW963138		
215285_s_at	1.41E-04	putative homeodomain transcription factor 1	AA927671	Hs.123637	NP_006599
215287_at	0.003171	Homo sapiens ELISC-1 mRNA, partial cds	AA975427	Hs.128434	
215313_x_at	0.007779	major histocompatibility complex, class I, C	AA573862	Hs.77961	NP_002107

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215318_at	0.028552	Consensus includes gb:AL049782.1 /DEF=Novel human gene mapping to chromosome 13. /FEA=mRNA /DB_XREF=gi:4902604 /UG=Hs.184938 Novel human gene mapping to chromosome 13	AL049782		
215336_at	0.005039	Consensus includes gb:AK002166.1 /DEF=Homo sapiens cDNA FLJ11304 fis, clone PLACE1009997, weakly similar to Rattus norvegicus A-kinase anchoring protein AKAP 220 mRNA. /FEA=mRNA /DB_XREF=gi:7023877 /UG=Hs.232076 A kinase (PRKA) anchor protein 11	AK002166		NP_652761
215338_s_at	1.26E-04	natural killer-tumor recognition sequence	AI688640	Hs.241493	NP_005376
215343_at	0.034221	Consensus includes gb:AF070587.1 /DEF=Homo sapiens clone :24741 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387960 /UG=Hs.25770 Homo sapiens clone 24741 mRNA sequence	AF070587		
215359_x_at	7.58E-04	GIOT-2 for gonadotropin inducible transcription repressor-2	AI758888	Hs.251371	
215366_at	0.041055	Consensus includes gb:AL353943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611). /FEA=mRNA /GEN=DKFZp761E0611 /PROD=hypothetical protein /DB_XREF=gi:7669982 /UG=Hs.284191 Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611)	AL353943		
215373_x_at	0.005039	Consensus includes gb:AK022213.1 /DEF=Homo sapiens cDNA FLJ12151 fis, clone MAMMA1000431. /FEA=mRNA /DB_XREF=gi:10433560 /UG=Hs.287493 hypothetical protein FLJ12151	AK022213		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215375_x_at	0.025245	Consensus includes gb:AK023938.1 /DEF=Homo sapiens cDNA FLJ13876 fis, clone THYRO1001401. /FEA=mRNA /DB_XREF=gi:10436033 /UG=Hs.287604 Homo sapiens cDNA FLJ13876 fis, clone THYRO1001401	AK023938		
215383_x_at	0.013223	Consensus includes gb:AL137312.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761K23121 (from clone DKFZp761K23121). /FEA=mRNA /DB_XREF=gi:6807797 /UG=Hs.293681 Homo sapiens mRNA; cDNA DKFZp761K23121 (from clone DKFZp761K23121)	AL137312		
215392_at	0.002418	AU148154 MAMMA1 Homo sapiens cDNA clone MAMMA1002744 3', mRNA sequence.	AU148154		
215399_s_at	1.26E-04	amplified in osteosarcoma	AI683900	Hs.76228	NP_006803
215404_x_at	0.016934	Consensus includes gb:AK024388.1 /DEF=Homo sapiens cDNA FLJ14326 fis, clone PLACE4000247. /FEA=mRNA /DB_XREF=gi:10436762 /UG=Hs.306678 Homo sapiens cDNA FLJ14326 fis, clone PLACE4000247	AK024388		
215424_s_at	5.03E-04	SKI-interacting protein	AV689564	Hs.79008	NP_036377
215434_x_at	0.0174	hypothetical protein DJ328E19.C1.1	AV684285	Hs.323463	NP_060410
215438_x_at	8.68E-04	G1 to S phase transition 1	BE906054	Hs.2707	NP_002085
215470_at	0.026891	Consensus includes gb:U21915.1 /DEF=Human chromosome 5q13.1 clone 5G8 mRNA. /FEA=mRNA /DB_XREF=gi:736412 /UG=Hs.14658 Human chromosome 5q13.1 clone 5G8 mRNA	U21915		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215492_x_at	0.040938	Consensus includes gb:AL035587 /DEF=Human DNA sequence from clone RP3-475N16 on chromosome 6p12.3-21.2. Contains the genes for CTG4A, pre-T cell receptor alpha, a novel protein similar to RPL7A (60S ribosomal protein L7A) and the 3 end of gene KIAA0240. Contains ESTs, STSs, GSSs an... /FEA=mRNA_2 /DB_XREF=gi:6002306 /UG=Hs.169002 Human pre TCR alpha mRNA, partial cds	AL035587		
215498_s_at	2.05E-04	mitogen-activated protein kinase kinase 3	AA780381	Hs.180533	NP_659732
215499_at	3.26E-04	mitogen-activated protein kinase kinase 3	AA780381	Hs.180533	NP_659732
215511_at	0.003171	Consensus includes gb:U19345.1 /DEF=Homo sapiens AR1 (TCF20) mRNA, partial cds. /FEA=mRNA /GEN=TCF20 /PROD=AR1 /DB_XREF=gi:2924754 /UG=Hs.201668 transcription factor 20 (AR1)	U19345		
215524_x_at	0.013088	immunoglobulin heavy constant mu	AW966434	Hs.300697	
215548_s_at	0.003171	Consensus includes gb:AB020724.1 /DEF=Homo sapiens mRNA for KIAA0917 protein, partial cds. /FEA=mRNA /GEN=KIAA0917 /PROD=KIAA0917 protein /DB_XREF=gi:4240322 /UG=Hs.27023 vesicle transport-related protein	AB020724		NP_057247
215549_x_at	0.00423	Consensus includes gb:AC005587 /DEF=Homo sapiens PAC clone RP5-988G15 from 7q33-q35 /FEA=CDS_2 /DB_XREF=gi:4156166 /UG=Hs.272525 Homo sapiens PAC clone RP5-988G15 from 7q33-q35	AC005587		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215566_x_at	0.013088	Consensus includes gb:AK024724.1 /DEF=Homo sapiens cDNA: FLJ21071 fis, clone CAS01789, highly similar to AF098668 Homo sapiens acyl-protein thioesterase mRNA. /FEA=mRNA /DB_XREF=gi:10437080 /UG=Hs.283655 lysophospholipase II	AK024724		NP_009191
215582_x_at	0.006603	Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 fis, clone MAMMA1001274. /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens cDNA FLJ12241 fis, clone MAMMA1001274	AK022303		
215588_x_at	0.003171	Consensus includes gb:AK024958.1 /DEF=Homo sapiens cDNA: FLJ21305 fis, clone COL02124. /FEA=mRNA /DB_XREF=gi:10437382 /UG=Hs.287658 Homo sapiens cDNA: FLJ21305 fis, clone COL02124	AK024958		
215599_at	0.040938	Consensus includes gb:X83300.1 /DEF=H.sapiens SMA4 mRNA. /FEA=mRNA /GEN=SMA4 /DB_XREF=gi:603028 /UG=Hs.289103 SMA4	X83300		
215600_x_at	0.005852	Consensus includes gb:AK022174.1 /DEF=Homo sapiens cDNA FLJ12112 fis, clone MAMMA1000043. /FEA=mRNA /DB_XREF=gi:10433510 /UG=Hs.288793 Homo sapiens cDNA FLJ12112 fis, clone MAMMA1000043	AK022174		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215602_at	6.68E-04	Consensus includes gb:AK024456.1 /DEF=Homo sapiens mRNA for FLJ00048 protein, partial cds. /FEA=mRNA /GEN=FLJ00048 /PROD=FLJ00048 protein /DB_XREF=gi:10440425 /UG=Hs.289034 Homo sapiens mRNA for FLJ00048 protein, partial cds	AK024456		
215603_x_at	8.94E-04	gamma-glutamyltransferase 1	AI344075	Hs.284380	
215604_x_at	0.013088	Consensus includes gb:AK023783.1 /DEF=Homo sapiens cDNA FLJ13721 fis, clone PLACE2000450. /FEA=mRNA /DB_XREF=gi:10435820 /UG=Hs.289035 Homo sapiens cDNA FLJ13721 fis, clone PLACE2000450	AK023783		
215608_at	0.033793	ESTs	AU144378	Hs.296650	
215623_x_at	0.005339	Consensus includes gb:AK002200.1 /DEF=Homo sapiens cDNA FLJ11338 fis, clone PLACE1010720, highly similar to Homo sapiens mRNA for chromosome-associated polypeptide-C. /FEA=mRNA /DB_XREF=gi:7023928 /UG=Hs.50758 SMC4 (structural maintenance of chromosomes 4, yeast)-like 1	AK002200		NP_005487
215628_x_at	0.001116	Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M193 (from clone DKFZp564M193). /FEA=mRNA /DB_XREF=gi:4500045 /UG=Hs.302053 Homo sapiens mRNA; cDNA DKFZp564M193 (from clone DKFZp564M193)	AL049285		
215629_s_at	0.013088	ESTs, Weakly similar to LEU2_HUMAN Leukemia associated protein 2 [H.sapiens]	AA905286	Hs.380846	
215633_x_at	0.041055	leukocyte specific transcript 1	AV713720	Hs.88411	
215646_s_at	3.26E-04	Homo sapiens versican Vint isoform, mRNA, partial cds	R94644	Hs.397468	
215667_x_at	0.016934	postmeiotic segregation increased 2-like 2	AI375694	Hs.177548	NP_777590

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215671_at	0.007779	AU144792 HEMBA1 Homo sapiens cDNA clone HEMBA1002973 3', mRNA sequence.	AU144792		
215684_s_at	5.07E-04	Consensus includes gb:AL096741.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586O0223 (from clone DKFZp586O0223). /FEA=mRNA /DB_XREF=gi:5419897 /UG=Hs.49005 hypothetical protein	AL096741		NP_115580
215694_at	0.018973	Consensus includes gb:AK022348.1 /DEF=Homo sapiens cDNA FLJ12286 fis, clone MAMMA1001768, weakly similar to CELL DIVISION CYCLE PROTEIN 48 HOMOLOG MJ1156. /FEA=mRNA /DB_XREF=gi:10433725 /UG=Hs.5555 hypothetical protein MGC5347	AK022348		NP_076968
215696_s_at	1.26E-04	Consensus includes gb:BC001404.1 /DEF=Homo sapiens, clone IMAGE:3139772, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3139772) /DB_XREF=gi:12655104 /UG=Hs.5716 KIAA0310 gene product	BC001404		
215698_at	3.26E-04	Consensus includes gb:AF007135.1 /DEF=Homo sapiens clone 23592 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852610 /UG=Hs.76272 retinoblastoma-binding protein 2	AF007135		
215716_s_at	0.033381	Consensus includes gb:L14561 /DEF=Homo sapiens plasma membrane calcium ATPase isoform 1 (ATP2B1) gene, alternative splice products, partial cds /FEA=mRNA_2 /DB_XREF=gi:4165324 /UG=Hs.78546 ATPase, Ca++ transporting, plasma membrane 1	L14561		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215719_x_at	0.033381	Consensus includes gb:X83493.1 /DEF=H.sapiens mRNA for FasApo-1 (clone pCRTM11-FasdeltaTM). /FEA=mRNA /DB_XREF=gi:971457 /UG=Hs.82359 tumor necrosis factor receptor superfamily, member 6	X83493		NP_690616
215735_s_at	0.002418	Consensus includes gb:AC005600 /DEF=Homo sapiens chromosome 16, P1 clone 109-9G (LANL), complete sequence /FEA=CDS_1 /DB_XREF=gi:3522919 /UG=Hs.90303 tuberous sclerosis 2	AC005600		
215739_s_at	6.68E-04	Consensus includes gb:AJ003062.1 /DEF=Homo sapiens mRNA for protein encoded by Saccharomyces cerevisiae SPC98 homologue. /FEA=mRNA /DB_XREF=gi:3152381 /UG=Hs.9884 spindle pole body protein	AJ003062		NP_006313
215749_s_at	0.009199	Consensus includes gb:AK001574.1 /DEF=Homo sapiens cDNA FLJ10712 fis, clone NT2RP3000919, highly similar to Rattus norvegicus golgi peripheral membrane protein p65 mRNA. /FEA=mRNA /DB_XREF=gi:7022910 /UG=Hs.4291 hypothetical protein FLJ23443	AK001574		NP_114105
215758_x_at	0.041055	HTF34-like ZNF gene; Homo sapiens chromosome 19, BAC 273239 (CIT-B-320G13), complete sequence.	AC007204		
215780_s_at	0.003539	Human DNA sequence from PAC 30P20 on chromosome Xq21.1-Xq21.3. Contains set pseudogene, ESTs and STS.	Z95126		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215806_x_at	0.026891	Consensus includes gb:M13231.1 /DEF=Human T-cell receptor aberrantly rearranged gamma-chain mRNA from cell line HPB-MLT. /FEA=mRNA /DB_XREF=gi:339168 /UG=Hs.274509 T cell receptor gamma constant 2	M13231		
215820_x_at	0.031948	Consensus includes gb:AK001861.1 /DEF=Homo sapiens cDNA FLJ10999 fis, clone PLACE1002529, highly similar to Homo sapiens mRNA for KIAA0713 protein. /FEA=mRNA /DB_XREF=gi:7023392 /UG=Hs.283881 KIAA0713 protein	AK001861		NP_055947
215832_x_at	3.26E-04	ubiquitin specific protease 2	AV722190	Hs.199160	NP_009097
215854_at	0.042466	ESTs	AU146050	Hs.287459	
215883_at	0.010205	Consensus includes gb:AK022326.1 /DEF=Homo sapiens cDNA FLJ12264 fis, clone MAMMA1001600. /FEA=mRNA /DB_XREF=gi:10433699 /UG=Hs.288481 Homo sapiens cDNA FLJ12264 fis, clone MAMMA1001600	AK022326		
215884_s_at	0.016934	Consensus includes gb:AK001029.1 /DEF=Homo sapiens cDNA FLJ10167 fis, clone HEMBA1003617, highly similar to Homo sapiens HRIHFB2157 mRNA. /FEA=mRNA /DB_XREF=gi:7022050 /UG=Hs.4552 ubiquilin 2	AK001029		NP_038472
215888_at	6.68E-04	Consensus includes gb:AK026889.1 /DEF=Homo sapiens cDNA: FLJ23236 fis, clone COL00725. /FEA=mRNA /DB_XREF=gi:10439854 /UG=Hs.288786 Homo sapiens cDNA: FLJ23236 fis, clone COL00725	AK026889		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215930_s_at	0.044231	Consensus includes gb:U73682.1 /DEF=Human meningioma-expressed antigen 11 (MEA11) mRNA, partial cds. /FEA=mRNA /GEN=MEA11 /PROD=meningioma-expressed antigen 11 /DB_XREF=gi:1916671 /UG=Hs.117242 meningioma expressed antigen 6 (coiled-coil proline-rich)	U73682		NP_005921
215933_s_at	2.05E-04	Consensus includes gb:Z21533.1 /DEF=H.sapiens HEX gene encoding homeobox related protein. /FEA=mRNA /GEN=HEX /PROD=homeobox related protein /DB_XREF=gi:32068 /UG=Hs.118651 hematopoietically expressed homeobox	Z21533		NP_002720
215938_s_at	0.022752	Consensus includes gb:AK001290.1 /DEF=Homo sapiens cDNA FLJ10428 fis, clone NT2RP1000376, highly similar to Homo sapiens mRNA; cDNA DKFZp434A102. /FEA=mRNA /DB_XREF=gi:7022453 /UG=Hs.120360 phospholipase A2, group VI (cytosolic, calcium-independent)	AK001290		
215952_s_at	0.041055	Consensus includes gb:AF090094.1 /DEF=Homo sapiens clone IMAGE 172979. /FEA=mRNA /DB_XREF=gi:4063629 /UG=Hs.125078 ornithine decarboxylase antizyme 1	AF090094		
215963_x_at	1.26E-04	Human DNA sequence from clone RP1-111B22 on chromosome 6q16-21, complete sequence.	Z98200		
215966_x_at	8.68E-04	glycerol kinase	AA292874	Hs.1466	NP_000158
215967_s_at	0.019657	lymphocyte antigen 9	AL582804	Hs.153042	NP_002339

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
215977_x_at	0.016934	Consensus includes gb:X68285.1 /DEF=H.sapiens mRNA for glycerol kinase. /FEA=mRNA /PROD=glycerol kinase /DB_XREF=gi:38413 /UG=Hs.1466 glycerol kinase	X68285		NP_000158
215978_x_at	0.001116	Consensus includes gb:AK021514.1 /DEF=Homo sapiens cDNA FLJ11452 fis, clone HEMBA1001435. /FEA=mRNA /DB_XREF=gi:10432710 /UG=Hs.148598 Homo sapiens cDNA FLJ11452 fis, clone HEMBA1001435	AK021514		
215983_s_at	0.023856	Consensus includes gb:D83768.1 /DEF=Human clone N9S Rep-8 mRNA, partial cds. /FEA=mRNA /GEN=Rep-8 /DB_XREF=gi:1913786 /UG=Hs.153678 reproduction 8	D83768		NP_005662
215990_s_at	0.003974	Consensus includes gb:S67779.1 /DEF=BCL5=Cys2-His2 zinc-finger transcription factor human, liver, mRNA, 2600 nt. /FEA=mRNA /DB_XREF=gi:459372 /UG=Hs.155024 B-cell CLLlymphoma 6 (zinc finger protein 51)	S67779		
216012_at	0.041055	Consensus includes gb:U43604.1 /DEF=Human unidentified mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:1171236 /UG=Hs.159901 Human unidentified mRNA, partial sequence	U43604		
216033_s_at	0.003171	Consensus includes gb:S74774.1 /DEF=p59fyn(T)=OKT3-induced calcium influx regulator human, Jurkat J6 T cell line, mRNA Partial, 1605 nt. /FEA=CDS /PROD=tyrosine kinase p59fyn(T) /DB_XREF=gi:802050 /UG=Hs.169370 FYN oncogene related to SRC, FGR, YES	S74774		NP_694593

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216035_x_at	5.64E-04	transcription factor 7-like 2 (T-cell specific, HMG-box)	AV721430	Hs.348412	NP_110383
216037_x_at	0.00228	transcription factor 7-like 2 (T-cell specific, HMG-box)	AA664011	Hs.348412	NP_110383
216041_x_at	0.041055	Consensus includes gb:AK023348.1 /DEF=Homo sapiens cDNA FLJ13286 fis, clone OVARC1001154, highly similar to Homo sapiens clone 24720 epithelin 1 and 2 mRNA. /FEA=mRNA /DB_XREF=gi:10435243 /UG=Hs.180577 granulin	AK023348		NP_002078
216054_x_at	5.03E-04	Human MLC1emb gene for embryonic myosin alkaline light chain, promoter and exon 1.	NM_002476		NP_002467
216063_at	0.001477	ESTs, Weakly similar to HBE_HUMAN HEMOGLOBIN EPSILON CHAIN [H.sapiens]	N55205	Hs.20205	
216071_x_at	0.010205	intelligence reducing insertion protein INGRIN; Homo sapiens OPA-containing protein (HOPA) gene, complete cds.	NM_005120		NP_005111
216088_s_at	0.00228	Consensus includes gb:AL078633 /DEF=Human DNA sequence from clone RP5-1005F21 on chromosome 20 Contains three isoforms of the gene PSMA7 (proteasome (prosome, macropain) subunit, alpha type, 7), the gene HRH3 encoding the Histamine H3 receptor, the gene KIAA0693 (similar to SSXT, sy... /FEA=CDS_4 /DB_XREF=gi:9408092 /UG=Hs.233952 proteasome (prosome, macropain) subunit, alpha type, 7	AL078633		
216090_x_at	0.023856	Consensus includes gb:AL078636.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929. /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929	AL078636		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216091_s_at	0.016934	Consensus includes gb:AF101784.1 /DEF=Homo sapiens b-TRCP variant E3RS-lkappaB mRNA, partial cds. /FEA=mRNA /PROD=b-TRCP variant E3RS-lkappaB /DB_XREF=gi:4165135 /UG=Hs.226434 beta-transducin repeat containing	AF101784		NP_378663
216095_x_at	0.00228	Consensus includes gb:AF057354.1 /DEF=Homo sapiens myotubularin-related protein 1a mRNA, partial cds. /FEA=mRNA /PROD=myotubularin-related protein 1a /DB_XREF=gi:5138901 /UG=Hs.23200 myotubularin related protein 1	AF057354		NP_789746
216100_s_at	6.79E-04	DKFZP586G011 protein	BG289527	Hs.234265	NP_056417
216105_x_at	0.03002	Consensus includes gb:X86428 /DEF=Homo sapiens PTPA gene for phosphotyrosyl phosphatase activator, exon 1 and joined CDS /FEA=mRNA_2 /DB_XREF=gi:974836 /UG=Hs.236963 protein phosphatase 2A, regulatory subunit B (PR 53)	X86428		
216110_x_at	0.046846	Consensus includes gb:AU147017 /FEA=EST /DB_XREF=gi:11008538 /DB_XREF=est:AU147017 /CLONE=HEMBB1002152 /UG=Hs.301905 Homo sapiens cDNA FLJ14080 fis, clone HEMBB1002152	AK024142		
216112_at	0.028893	AU157200 PLACE1 Homo sapiens cDNA clone PLACE1006617 3', mRNA sequence.	AU157200		
216125_s_at	0.001323	Consensus includes gb:AF064606.1 /DEF=Homo sapiens KB07 protein mRNA, partial cds. /FEA=mRNA /PROD=KB07 protein /DB_XREF=gi:3152665 /UG=Hs.306242 Homo sapiens KB07 protein mRNA, partial cds	AF064606		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216126_at	0.022752	Consensus includes gb:AK022418.1 /DEF=Homo sapiens cDNA FLJ12356 fis, clone MAMMA1002339. /FEA=mRNA /DB_XREF=gi:10433811 /UG=Hs.306635 Homo sapiens cDNA FLJ12356 fis, clone MAMMA1002339	AK022418		
216187_x_at	0.013223	Consensus includes gb:AF222691.1 /DEF=Homo sapiens Alu repeat (LNX1) mRNA sequence. /FEA=mRNA /DB_XREF=gi:12655850 /UG=Hs.307008 Homo sapiens Alu repeat (LNX1) mRNA sequence	AF222691		
216191_s_at	0.004341	Consensus includes gb:X72501.1 /DEF=H.sapiens mRNA for soluble delta TCR. /FEA=CDS /PROD=gamma-delta T-cell receptor /DB_XREF=gi:298104 /UG=Hs.307187 H.sapiens mRNA for soluble delta TCR	X72501		
216202_s_at	0.016686	Consensus includes gb:U15555.1 /DEF=Human serine palmitoyltransferase (LCB2) mRNA, partial cds. /FEA=mRNA /GEN=LCB2 /PROD=serine palmitoyltransferase /DB_XREF=gi:1001944 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2	U15555		NP_004854
216207_x_at	0.045316	immunoglobulin kappa constant	AW408194	Hs.306357	
216218_s_at	1.26E-04	Consensus includes gb:AK023546.1 /DEF=Homo sapiens cDNA FLJ13484 fis, clone PLACE1003888, weakly similar to 1-PHOSPHATIDYLINOSITOL-4,5-BISPHOSPHATE PHOSPHODIESTERASE DELTA 1 (EC 3.1.4.11). /FEA=mRNA /DB_XREF=gi:10435512 /UG=Hs.54886 phospholipase C, epsilon 2	AK023546		NP_055999

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216221_s_at	7.58E-04	Consensus includes gb:D87078.2 /DEF=Homo sapiens mRNA for KIAA0235 protein, partial cds. /FEA=mRNA /GEN=KIAA0235 /PROD=KIAA0235 protein /DB_XREF=gi:6634004 /UG=Hs.6151 pumilio (Drosophila) homolog 2	D87078		NP_056132
216266_s_at	0.026891	Consensus includes gb:AK025637.1 /DEF=Homo sapiens cDNA: FLJ21984 fis, clone HEP06222, highly similar to AF111162 Homo sapiens guanine nucleotide exchange factor mRNA. /FEA=mRNA /DB_XREF=gi:10438216 /UG=Hs.94631 brefeldin A-inhibited guanine nucleotide-exchange protein 1	AK025637		NP_006412
216268_s_at	1.28E-04	Consensus includes gb:U77914.1 /DEF=Human soluble protein Jagged mRNA, partial cds. /FEA=mRNA /PROD=soluble protein Jagged /DB_XREF=gi:1684889 /UG=Hs.91143 jagged 1 (Alagille syndrome)	U77914		NP_000205
216274_s_at	3.26E-04	signal peptidase complex (18kD)	N99438	Hs.9534	NP_055115
216282_x_at	0.016934	Consensus includes gb:AJ224143.1 /DEF=Homo sapiens mRNA variant alfa for RNA polymerase II subunit 3. /FEA=mRNA /GEN=RPB3 /PROD=RNA polymerase II subunit /DB_XREF=gi:2920708 /UG=Hs.79402 polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AJ224143		NP_116558
216288_at	6.68E-04	AU159276 THYRO1 Homo sapiens cDNA clone THYRO1001262 3', mRNA sequence.	AU159276		NP_006630
216295_s_at	0.010205	Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene	X81636		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216296_at	0.022988	Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene	X81636		
216304_x_at	0.041055	Homo sapiens FTSH gene for putative ATPases, exons 1 and 2 and join CDS.	AJ295618		
216308_x_at	0.007779	Consensus includes gb:AK026752.1 /DEF=Homo sapiens cDNA: FLJ23099 fis, clone LNG07460, highly similar to AF113251 Homo sapiens putative 2-hydroxyacid dehydrogenase mRNA. /FEA=mRNA /DB_XREF=gi:10439675 /UG=Hs.155742 glyoxylate reductasehydroxypyruvate reductase	AK026752		NP_036335
216321_s_at	0.00161	Consensus includes gb:X03348.1 /DEF=Human mRNA for beta-glucocorticoid receptor (clone OB10). /FEA=mRNA /PROD=beta-glucocorticoid receptor /DB_XREF=gi:31681 /UG=Hs.75772 nuclear receptor subfamily 3, group C, member 1	X03348		NP_000167
216323_x_at	0.001303	alpha-tubulin isotype H2-alpha; Human alpha-tubulin isotype H2-alpha gene, last exon.	K03460		
216379_x_at	3.26E-04	Consensus includes gb:AK000168.1 /DEF=Homo sapiens cDNA FLJ20161 fis, clone COL09252, highly similar to L33930 Homo sapiens CD24 signal transducer mRNA. /FEA=mRNA /DB_XREF=gi:7020079 /UG=Hs.332045 Homo sapiens cDNA FLJ20161 fis, clone COL09252, highly similar to L33930 Homo sapiens CD24 signal transducer mRNA	AK000168		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216383_at	2.05E-04	Homo sapiens chromosome X clone Qc-7G6, QLL-F1720, QLL-C1335, Qc-8B7, Qc-11H12, Qc-7F6, QLL-E153, Qc-10E8, Qc-10B7 map q28, complete sequence.	U52111		
216388_s_at	0.00884	R2; Human putative G-protein-coupled receptor (GPR16) gene, complete cds.	U33448		
216389_s_at	0.021312	Consensus includes gb:AF283773.2 /DEF=Homo Sapiens clone TCBAP0888 mRNA sequence. /FEA=mRNA /DB_XREF=gi:10281743 /UG=Hs.283976 hypthetical protein PRO2389	AF283773		NP_079506
216409_at	0.001116	Consensus includes gb:AL390168.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547D104 (from clone DKFZp547D104). /FEA=mRNA /GEN=DKFZp547D104 /PROD=hypothetical protein /DB_XREF=gi:9368863 /UG=Hs.287371 hypothetical protein DKFZp547D104	AL390168		
216443_at	0.034317	Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fis, clone ADSh00804. /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens cDNA: FLJ20962 fis, clone ADSh00804	AK024615		
216457_s_at	3.26E-04	Consensus includes gb:AK026080.1 /DEF=Homo sapiens cDNA: FLJ22427 fis, clone HRC09013. /FEA=mRNA /DB_XREF=gi:10438814 /UG=Hs.288883 splicing factor 3a, subunit 1, 120kD	AK026080		
216483_s_at	0.020793	Hypothetical partial human protein; Homo sapiens chromosome 19, cosmid R33729, complete sequence.	AC005339		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216484_x_at	1.26E-04	Consensus includes gb:L24521.1 /DEF=Human transformation-related protein mRNA, 3 end. /FEA=mRNA /PROD=transformation-related protein /DB_XREF=gi:403459 /UG=Hs.300705 Human transformation-related protein mRNA, 3 end	L24521		NP_004485
216508_x_at	0.016934	similar to nonhistone chromosomal protein HMG-1 [Homo sapiens]; probable pseudogene; similar to P09429 (PID:g123369); H_NH0244E06.1; Homo sapiens BAC clone RP11-244E6 from 2, complete sequence.	AC007277		
216511_s_at	0.005852	Consensus includes gb:AJ270770 /DEF=Homo sapiens partial TCF-4 gene for T-cell transcription factor-4, exon 1 and joined CDS features /FEA=mRNA_1 /DB_XREF=gi:9188625 /UG=Hs.283857 Homo sapiens partial TCF-4 gene for T-cell transcription factor-4, exon 1 and joined CDS features	AJ270770		
216520_s_at	0.005852	Homo sapiens HDCMB21P gene, complete cds.	AF072098		
216526_x_at	0.041055	Consensus includes gb:AK024836.1 /DEF=Homo sapiens cDNA: FLJ21183 fis, clone CAS11634, highly similar to HSHLACW07 Homo sapiens mRNA for human leukocyte antigen C alpha chain. /FEA=mRNA /DB_XREF=gi:10437242 /UG=Hs.277477 major histocompatibility complex, class I, C	AK024836		NP_002108

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216550_x_at	0.026891	Consensus includes gb:X80821.1 /DEF=H.sapiens mRNA for ribosomal protein L18a homologue. /FEA=mRNA /PROD=ribosomal protein L18a homologue /DB_XREF=gi:527579 /UG=Hs.302177 H.sapiens mRNA for ribosomal protein L18a homologue	X80821		NP_056023
216556_x_at	0.021197	Consensus includes gb:AL135926 /DEF=Human DNA sequence from clone RP11-375F2 on chromosome 1 Contains a pseudogene similar to UBL1 (ubiquitin-like 1 (sentrin)), a pseudogene similar to ribosomal protein L29, ESTs, STSs and GSSs /FEA=CDS_1 /DB_XREF=gi:9801286 /UG=Hs.302113 Human DNA sequence from clone RP11-375F2 on chromosome 1 Contains a pseudogene similar to UBL1 (ubiquitin-like 1 (sentrin)), a pseudogene similar to ribosomal protein L29, ESTs, STSs and GSSs	AL135926		
216563_at	6.68E-04	Consensus includes gb:X80821.1 /DEF=H.sapiens mRNA for ribosomal protein L18a homologue. /FEA=mRNA /PROD=ribosomal protein L18a homologue /DB_XREF=gi:527579 /UG=Hs.302177 H.sapiens mRNA for ribosomal protein L18a homologue	X80821		NP_056023

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216565_x_at	0.016934	dJ781L3.1 (similar to IFITM3 (interferon induced transmembrane protein 3 (1-8U))) match: proteins: Sw:Q01628 Sw:Q01629 Tr:Q9R175 Tr:Q9R176 Sw:Q91499 Sw:P26376 Tr:O88728 Sw:P13164; Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon induced transmembrane protein 3 (1-8U)), STSs and GSSs, complete sequence.	AL121994		
216570_x_at	0.004341	Consensus includes gb:AL096829 /DEF=Human DNA sequence from clone RP4-595K12 on chromosome 1p31.2-31.3 Contains a pseudogene similar to 60S RPL29 (ribosomal protein L29 (cell surface heparin binding protein HIP)), a chromosome 1 specific mRNA (KIAA0499), a novel mRNA (KIAA0433), EST... /FEA=CDS /DB_XREF=gi:6634461 /UG=Hs.302120 Human DNA sequence from clone RP4-595K12 on chromosome 1p31.2-31.3 Contains a pseudogene similar to 60S RPL29 (ribosomal protein L29 (cell surface heparin binding protein HIP)), a chromosome 1 specific mRNA (KIAA0499), a novel mRNA (KIAA0433), ESTs, STSs,	AL096829		
216593_s_at	0.003171	pseudogene of PIGC; see also AB000360 putative; Homo sapiens PIGCP1 pseudogene.	AB000359		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216606_x_at	0.029453	Consensus includes gb:AL050332 /DEF=Human DNA sequence from clone RP4-570F3 on chromosome 6 Contains a gene similar to Rattus norvegicus synaptic ras GTPase-activating protein p135, the CICK0721Q.5 (polypeptide from patented cDNA Em:E06811) gene, the PHF1 (PHD finger protein 1) gen... /FEA=mRNA_1 /DB_XREF=gi:6010176 /UG=Hs.306238 acyl-protein	AL050332		
216640_s_at	0.041055	Consensus includes gb:AK026926.1 /DEF=Homo sapiens cDNA: FLJ23273 fis, clone HEP02611, highly similar to HSU79278 Human protein disulfide isomerase-related protein P5 mRNA. /FEA=mRNA /DB_XREF=gi:10439898 /UG=Hs.182429 protein disulfide isomerase-related protein	AK026926		
216652_s_at	1.26E-04	Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872). /FEA=mRNA /DB_XREF=gi:6807841 /UG=Hs.306454 Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872)	AL137673		
216739_at	0.009716	Consensus includes gb:AK024527.1 /DEF=Homo sapiens cDNA: FLJ20874 fis, clone ADKA02818. /FEA=mRNA /DB_XREF=gi:10436829 /UG=Hs.306684 Homo sapiens cDNA: FLJ20874 fis, clone ADKA02818	AK024527		
216751_at	0.02145	Consensus includes gb:AK024879.1 /DEF=Homo sapiens cDNA: FLJ21226 fis, clone COL00721. /FEA=mRNA /DB_XREF=gi:10437291 /UG=Hs.306715 Homo sapiens cDNA: FLJ21226 fis, clone COL00721	AK024879		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216756_at	8.68E-04	Consensus includes gb:AK024995.1 /DEF=Homo sapiens cDNA: FLJ21342 fis, clone COL02673. /FEA=mRNA /DB_XREF=gi:10437426 /UG=Hs.306739 Homo sapiens cDNA: FLJ21342 fis, clone COL02673	AK024995		
216791_at	0.006096	Consensus includes gb:AK026449.1 /DEF=Homo sapiens cDNA: FLJ22796 fis, clone KAIA2544. /FEA=mRNA /DB_XREF=gi:10439318 /UG=Hs.306850 Homo sapiens cDNA: FLJ22796 fis, clone KAIA2544	AK026449		
216804_s_at	0.011725	Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fis, clone LNG10773. /FEA=mRNA /DB_XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cDNA: FLJ23564 fis, clone LNG10773	AK027217		
216833_x_at	0.034317	Consensus includes gb:U05255.1 /DEF=Human glycoporphin HeP2 mRNA, partial cds. /FEA=mRNA /PROD=glycoporphin HeP2 /DB_XREF=gi:454085 /UG=Hs.307185 Human glycoporphin HeP2 mRNA, partial cds	U05255		NP_002091
216835_s_at	2.05E-04	Consensus includes gb:AF035299.1 /DEF=Homo sapiens clone 23863 mRNA, partial cds. /FEA=mRNA /DB_XREF=gi:2661055 /UG=Hs.103854 docking protein 1, 62kD (downstream of tyrosine kinase 1)	AF035299		NP_001372
216841_s_at	7.58E-04	Consensus includes gb:X15132.1 /DEF=Human mRNA for manganese containing superoxide dismutase (EC 1.15.1.1). /FEA=mRNA /DB_XREF=gi:34794 /UG=Hs.318885 superoxide dismutase 2, mitochondrial	X15132		NP_000627

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216858_x_at	0.018973	Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.332731 Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722)	AL080112		
216870_x_at	0.007779	Consensus includes gb:AF264787.1 /DEF=Homo sapiens BCMS-upstream neighbor (BCMSUN) mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:11141510 /UG=Hs.43628 deleted in lymphocytic leukemia, 2	AF264787		NP_006012
216899_s_at	0.007779	Homo sapiens PAC clone RP5-1139P1 from 7p11.2-p21, complete sequence.	AC003999		
216902_s_at	0.009716	Consensus includes gb:AF001549 /DEF=Human Chromosome 16 BAC clone CIT987SK-A-270G1 /FEA=mRNA_1 /DB_XREF=gi:3355302 /UG=Hs.110103 RNA polymerase I transcription factor RRN3	AF001549		
216903_s_at	1.32E-04	Consensus includes gb:AK022697.1 /DEF=Homo sapiens cDNA FLJ12635 fis, clone NT2RM4001865, highly similar to Homo sapiens mRNA for atopy related autoantigen CALC. /FEA=mRNA /DB_XREF=gi:10434244 /UG=Hs.61628 calcium binding atopy-related autoantigen 1	AK022697		NP_006068
216942_s_at	1.26E-04	Consensus includes gb:D28586.1 /DEF=Human mRNA for LFA-3(delta D2), partial cds. /FEA=mRNA /PROD=LFA-3(delta D2) /DB_XREF=gi:466540 /UG=Hs.75626 CD58 antigen, (lymphocyte function-associated antigen 3)	D28586		NP_001770

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216950_s_at	0.003171	Consensus includes gb:X14355.1 /DEF=Human mRNA for high affinity Fc receptor (FcRI) b form. /FEA=mRNA /DB_XREF=gi:31333 /UG=Hs.77424 Fc fragment of IgG, high affinity Ia, receptor for (CD64)	X14355		
216988_s_at	2.05E-04	Homo sapiens (clone hh18) protein tyrosine phosphatase (ptp-IV1r) gene, 5' end of cds.	L48722		
216993_s_at	0.016686	Consensus includes gb:U32169 /DEF=Human pro-a2 chain of collagen type XI (COL11A2) gene, complete cds /FEA=mRNA_3 /DB_XREF=gi:1000744 /UG=Hs.121509 collagen, type XI, alpha 2	U32169		
217003_s_at	0.041795	Consensus includes gb:AJ132823.1 /DEF=Homo sapiens mRNA for tMDC II, isoform d. /FEA=mRNA /DB_XREF=gi:4468348 /UG=Hs.194768 a disintegrin and metalloproteinase domain 5	AJ132823		
217009_at	0.018443	Consensus includes gb:AL121974 /DEF=Human DNA sequence from clone RP3-417L20 on chromosome 6p12-21.3. Contains the 5' end of the gene for cysteine-rich secretory protein 3 (CRISP-3, SGP28), the PGK2 gene for phosphoglycerate kinase 2, STSs and GSSs /FEA=CDS /DB_XREF=gi:9828129 /UG=Hs.78771 phosphoglycerate kinase 1	AL121974		
217016_x_at	0.003171	Consensus includes gb:AK026825.1 /DEF=Homo sapiens cDNA: FLJ23172 fis, clone LNG10005. /FEA=mRNA /DB_XREF=gi:10439771 /UG=Hs.306885 Homo sapiens cDNA: FLJ23172 fis, clone LNG10005	AK026825		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217041_at	0.017634	Consensus includes gb:AL162057.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N0524 (from clone DKFZp761N0524). /FEA=mRNA /DB_XREF=gi:7328105 /UG=Hs.91622 neuronal pentraxin receptor	AL162057		NP_478058
217042_at	0.003171	Consensus includes gb:AL096716.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M1462 (from clone DKFZp564M1462); partial cds. /FEA=mRNA /GEN=DKFZp564M1462 /PROD=hypothetical protein /DB_XREF=gi:5419850 /UG=Hs.226007 DKFZP564M1462 protein	AL096716		
217043_s_at	1.26E-04	Consensus includes gb:U95822.1 /DEF=Human putative transmembrane GTPase mRNA, partial cds. /FEA=mRNA /GEN=fzo /PROD=putative transmembrane GTPase /DB_XREF=gi:2252803 /UG=Hs.197877 hypothetical protein FLJ20693	U95822		
217078_s_at	0.003597	Consensus includes gb:AJ010102.1 /DEF=Homo Sapiens mRNA for Natural killer cell IRC1c gene. /FEA=mRNA /GEN=IRC1c /DB_XREF=gi:5706457 /UG=Hs.9688 leukocyte membrane antigen	AJ010102		
217118_s_at	2.05E-04	Consensus includes gb:AK025608.1 /DEF=Homo sapiens cDNA: FLJ21955 fis, clone HEP05265. /FEA=mRNA /DB_XREF=gi:10438177 /UG=Hs.13255 KIAA0930 protein	AK025608		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217122_s_at	0.033381	Consensus includes gb:AL031282 /DEF=Human DNA sequence from clone 283E3 on chromosome 1p36.21-36.33. Contains the alternatively spliced gene for Matrix Metalloproteinase in the Female Reproductive tract MIFR1, -2, MMP2122A, -B and -C, a novel gene, the alternatively spliced CDC2L2 ... /FEA=mRNA_6 /DB_XREF=gi:3860395 /UG=Hs.214646 KIAA0447 gene product	AL031282		
217125_at	0.005692	Homo sapiens UBBP2 pseudogene for ubiquitin UBB.	X04802		
217143_s_at	0.02145	Consensus includes gb:X06557.1 /DEF=Human mRNA for TCR-delta chain. /FEA=mRNA /DB_XREF=gi:37003 /UG=Hs.2014 T cell receptor delta locus	X06557		
217144_at	1.28E-04	Homo sapiens UBBP1 pseudogene for ubiquitin UBB.	X04801		
217164_at	0.010205	Consensus includes gb:AK024108.1 /DEF=Homo sapiens cDNA FLJ14046 fis, clone HEMBA1006461. /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapiens cDNA FLJ14046 fis, clone HEMBA1006461	AK024108		
217167_x_at	0.003862	Consensus includes gb:AJ252550 /DEF=Homo sapiens partial GK gene for glycerol kinase, exon 1 (glycerol kinase deficiency case) /FEA=CDS_1 /DB_XREF=gi:5834425 /UG=Hs.1466 glycerol kinase	AJ252550		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217168_s_at	0.026891	Consensus includes gb:AF217990.1 /DEF=Homo sapiens clone PP1722 unknown mRNA. /FEA=mRNA /PROD=unknown /DB_XREF=gi:10441910 /UG=Hs.146393 homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	AF217990		NP_055500
217200_x_at	7.58E-04	Consensus includes gb:U06715.1 /DEF=Human cytochrome B561, HCYTO B561, mRNA, partial cds. /FEA=mRNA /GEN=B561 /PROD=HCYTO B561 /DB_XREF=gi:476590 /UG=Hs.153028 cytochrome b-561	U06715		
217208_s_at	0.020576	match: proteins: Tr:Q62402 Sw:Q62696 Sw:Q12959; Human DNA sequence from clone RP5-1061C18 on chromosome 3p36.22-36.33 Contains the 5' part of the DLG1 gene encoding the discs protein (large Drosophila homolog 1, presynaptic protein SAP97) and a CpG island, complete sequence.	AL121981		
217209_at	0.005673	Human gene for carcinoembryonic antigen subdomains A and B.	X16454		
217211_at	6.79E-04	Human beta-cytoplasmic actin (ACTBP9) pseudogene.	D50604		
217212_s_at	7.44E-04	Human DNA sequence from phage LAW2 from a contig from the tip of the short arm of chromosome 16, spanning 2Mb of 16p13.3 Contains Interleukin 9 receptor pseudogene.	Z84723		
217226_s_at	0.026891	Consensus includes gb:M95929.1 /DEF=Human homeobox protein (PHOX1) mRNA, 3 end. /FEA=mRNA /GEN=PHOX1 /PROD=homeobox protein /DB_XREF=gi:189946 /UG=Hs.155606 paired mesoderm homeo box 1	M95929		
217232_x_at	0.001116	Homo sapiens mutant beta-globin (HBB) gene, complete cds.	AF059180		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217249_x_at	0.005852	cytochrome C oxidase; match to P14406 (PID:g117121); H_RG162B04.1; Homo sapiens BAC clone CTB-162B4 from 4, complete sequence.	AC004544		
217274_x_at	0.001205	Consensus includes gb:X52005.1 /DEF=H.sapiens skeletal embryonic myosin light chain 1 (MLC1) mRNA. /FEA=mRNA /GEN=MCL1 /PROD=myosin light chain 1 /DB_XREF=gi:34677 /UG=Hs.159218 H.sapiens skeletal embryonic myosin light chain 1 (MLC1) mRNA	X52005		
217286_s_at	0.007779	Consensus includes gb:BC001805.1 /DEF=Homo sapiens, clone IMAGE:3543670, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3543670) /DB_XREF=gi:12804742 /UG=Hs.240615 hypothetical protein FLJ13556 similar to N-myc downstream regulated 3	BC001805		
217294_s_at	0.029453	Consensus includes gb:U88968.1 /DEF=Human alpha enolase like 1 (ENO1L1) mRNA, partial cds. /FEA=mRNA /GEN=ENO1L1 /PROD=alpha enolase like 1 /DB_XREF=gi:3282242 /UG=Hs.254105 enolase 1, (alpha)	U88968		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217356_s_at	0.033381	Consensus includes gb:S81916.1 /DEF=phosphoglycerate kinase {alternatively spliced} human, phosphoglycerate kinase deficient patient with episodes of muscl, mRNA Partial Mutant, 307 nt. /FEA=mRNA /PROD=phosphoglycerate kinase /DB_XREF=gi:1470308 /UG=Hs.169313 Phosphoglycerate kinase {alternatively spliced} human, phosphoglycerate kinase deficient patient with episodes of muscl, mRNA Partial Mutant, 307 nt	S81916		
217362_x_at	0.001551	Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mRNA /DB_XREF=gi:5915893 /UG=Hs.167385 Homo sapiens MHC class II antigen HLA-DRB6 mRNA, partial cds	AF005487		
217371_s_at	0.010644	Consensus includes gb:Y09908.1 /DEF=H.sapiens mRNA for interleukin-15. /FEA=mRNA /GEN=IL-15 /PROD=interleukin-15 /DB_XREF=gi:2143255 /UG=Hs.168132 interleukin 15	Y09908		
217393_x_at	0.021165	match: proteins: Sw:P52490 Sw:P35128 Sw:Q29503 Sw:P52492 Tr:Q42045 Tr:Q9ZVA6 Tr:Q9ZTZ8 Tr:Q42897 Sw:P35131 Sw:P35132 Sw:P35133 Sw:P35134 Sw:P35135 Wp:CE19850 Tr:O13685 Tr:Q43780 Sw:Q16781 Tr:O94098; Human DNA sequence from clone RP3-526F5 on chromosome Xq26.3-28. Contains the gene for a novel protein similar to UBE2N (ubiquitin-conjugating enzyme E2N (homologous to yeast UBC13)), ESTs, STSs and GSSs, complete sequence.	AL109622		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217394_at	0.026891	Consensus includes gb:AE000659 /DEF=Homo sapiens T-cell receptor alpha delta locus from bases 250472 to 501670 (section 2 of 5) of the Complete Nucleotide Sequence /FEA=CDS_9 /DB_XREF=gi:2358025 /UG=Hs.272547 Homo sapiens T-cell receptor alpha delta locus from bases 250472 to 501670 (section 2 of 5) of the Complete Nucleotide Sequence	AE000659		
217403_s_at	0.041795	Consensus includes gb:AC074331 /DEF=Homo sapiens chromosome 19, BAC CIT-HSPC_204F22 (BC228680), complete sequence; contains bacterial insertion element /FEA=CDS_4 /DB_XREF=gi:9502399 /UG=Hs.9615 myosin regulatory light chain 2, smooth muscle isoform	AC074331		
217414_x_at	3.56E-04	Human alpha-globin gene with flanks.	NM_000517		NP_000508
217418_x_at	0.018693	Consensus includes gb:X12530.1 /DEF=Human mRNA for B lymphocyte antigen CD20 (B1, Bp35). /FEA=mRNA /DB_XREF=gi:29773 /UG=Hs.89751 membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide)	X12530		NP_690606
217427_s_at	4.42E-04	Consensus includes gb:X75296.1 /DEF=H.sapiens TUP1-like enhancer of split gene 1 (TUPLE1) mRNA. /FEA=mRNA /GEN=TUPLE1 /PROD=TUP1 like enhancer of SPLIT gene 1 /DB_XREF=gi:434982 /UG=Hs.172350 HIR (histone cell cycle regulation defective, S. cerevisiae) homolog A	X75296		NP_003316

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217437_s_at	2.05E-04	Consensus includes gb:AB029026.1 /DEF=Homo sapiens mRNA for KIAA1103 protein, partial cds. /FEA=mRNA /GEN=KIAA1103 /PROD=KIAA1103 protein /DB_XREF=gi:5689542 /UG=Hs.173159 transforming, acidic coiled-coil containing protein 1	AB029026		NP_006274
217456_x_at	0.00161	Consensus includes gb:M31183.1 /DEF=Homo sapiens (cosmid clone cd3.3) MHC class I HLA-B51 mRNA, 3 end of cds. /FEA=mRNA /GEN=HLA-B /PROD=MHC class I lymphocyte antigen /DB_XREF=gi:601851 /UG=Hs.181392 major histocompatibility complex, class I, E	M31183		NP_005507
217457_s_at	0.028552	Consensus includes gb:X63465.1 /DEF=H.sapiens hGDS mRNA for smg GDS. /FEA=CDS /GEN=hGDS /PROD=smg GDS /DB_XREF=gi:32079 /UG=Hs.7940 RAP1, GTP-GDP dissociation stimulator 1	X63465		NP_066982
217475_s_at	0.028999	Consensus includes gb:AC002073 /DEF=Human PAC clone RP3-515N1 from 22q11.2-q22 /FEA=CDS_1 /DB_XREF=gi:2078469 /UG=Hs.278027 LIM domain kinase 2	AC002073		
217479_at	0.034317	Consensus includes gb:AL110201.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622). /FEA=mRNA /DB_XREF=gi:5817120 /UG=Hs.278435 Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622)	AL110201		
217480_x_at	0.005411	Ig kappa chain; Human kappa-immunoglobulin germline pseudogene (cos118) variable region (subgroup V kappa I).	M20812		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217497_at	0.016934	ESTs, Weakly similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	AW613387	Hs.388345	NP_001944
217499_x_at	2.17E-04	olfactory receptor, family 7, subfamily E, member 38 pseudogene	AW874308	Hs.409186	
217502_at	0.004341	interferon-induced protein with tetratricopeptide repeats 2	BE888744	Hs.169274	NP_001538
217512_at	0.033491	ESTs, Highly similar to 1312232A kininogen L,high MW [Homo sapiens] [H.sapiens]	BG398937	Hs.289056	NP_000884
217529_at	0.04734	ESTs, Moderately similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	BE547674	Hs.129568	
217534_at	0.015516	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AV720514	Hs.282721	
217540_at	1.26E-04	ESTs, Moderately similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	AA721025	Hs.293253	
217552_x_at	3.26E-04	th43e02.x1 NCI_CGAP_Lym12 Homo sapiens cDNA clone IMAGE:2121050 3' similar to contains Alu repetitive element; mRNA sequence.	AI432713		
217560_at	0.02008	ESTs, Weakly similar to hypothetical protein FLJ11267 [Homo sapiens] [H.sapiens]	AI879383	Hs.370291	
217580_x_at	0.041795	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW301806	Hs.150551	
217610_at	0.042057	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AL047879	Hs.293865	
217614_at	0.02145	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AI439416	Hs.165728	
217627_at	7.58E-04	hypothetical protein FLJ30921	BE515346	Hs.278871	NP_689573
217629_at	0.00228	ESTs, Moderately similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	AA365670	Hs.261526	
217673_x_at	0.013223	ESTs, Highly similar to GBAS_HUMAN Guanine nucleotide binding protein G(S), alpha subunit (Adenylate cyclase-stimulating G alpha protein) [H.sapiens]	AA650558	Hs.394484	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217679_x_at	0.005852	tx67h02.x1 NCI_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2274675 3' similar to contains Alu repetitive element;, mRNA sequence.	AI683552		
217682_at	0.004341	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW503390	Hs.392578	
217695_x_at	0.007934	ESTs, Weakly similar to I57588 HSrel-1 - human (fragment) [H.sapiens]	BF942161	Hs.343402	
217707_x_at	3.26E-04	ESTs	AI535683	Hs.405576	
217713_x_at	0.005039	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AA126763	Hs.182296	
217715_x_at	0.010205	ESTs, Weakly similar to hypothetical protein FLJ20294 [Homo sapiens] [H.sapiens]	BE045142	Hs.274907	
217720_at	0.02145	gb:NM_016139.1 /DEF=Homo sapiens 16.7Kd protein (LOC51142), mRNA. /FEA=mRNA /GEN=LOC51142 /PROD=16.7Kd protein /DB_XREF=gi:7705850 /UG=Hs.180859 16.7Kd protein /FL=gb:BC003079.1 gb:AF078845.1 gb:NM_016139.1	NM_016139		NP_057223
217724_at	0.041055	Consensus includes gb:AF131807.1 /DEF=Homo sapiens clone 25076 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4406639 /UG=Hs.165998 PAI-1 mRNA-binding protein /FL=gb:AL080119.1 gb:NM_015640.1	AF131807		NP_056455
217725_x_at	0.02145	gb:NM_015640.1 /DEF=Homo sapiens PAI-1 mRNA-binding protein (PAI-RBP1), mRNA. /FEA=mRNA /GEN=PAI-RBP1 /PROD=PAI-1 mRNA-binding protein /DB_XREF=gi:7661625 /UG=Hs.165998 PAI-1 mRNA-binding protein /FL=gb:AL080119.1 gb:NM_015640.1	NM_015640		NP_056455

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217727_x_at	1.26E-04	gb:NM_018206.1 /DEF=Homo sapiens vacuolar protein sorting 35 (yeast homolog) (VPS35), mRNA. /FEA=mRNA /GEN=VPS35 /PROD=vacuolar protein sorting 35 (yeast homolog) /DB_XREF=gi:8922645 /UG=Hs.264190 vacuolar protein sorting 35 (yeast homolog) /FL=gb:AF186382.1 gb:AL136888.1 gb:BC002414.1 gb:AF191298.2 gb:NM_018206.1 gb:AF175265.1 gb:AF183418.1	NM_018206		NP_060676
217728_at	3.26E-04	gb:NM_014624.2 /DEF=Homo sapiens S100 calcium-binding protein A6 (calcyclin) (S100A6), mRNA. /FEA=mRNA /GEN=S100A6 /PROD=S100 calcium-binding protein A6 /DB_XREF=gi:9845517 /UG=Hs.275243 S100 calcium-binding protein A6 (calcyclin) /FL=gb:BC001431.1 gb:NM_014624.2	NM_014624		NP_055439
217729_s_at	6.68E-04	gb:NM_001130.3 /DEF=Homo sapiens amino-terminal enhancer of split (AES), mRNA. /FEA=mRNA /GEN=AES /PROD=amino-terminal enhancer of split /DB_XREF=gi:6382065 /UG=Hs.244 amino-terminal enhancer of split /FL=gb:AF072902.1 gb:U04241.1 gb:NM_001130.3	NM_001130		NP_001121
217730_at	0.005852	gb:NM_022152.1 /DEF=Homo sapiens PP1201 protein (PP1201), mRNA. /FEA=mRNA /GEN=PP1201 /PROD=PP1201 protein /DB_XREF=gi:11545897 /UG=Hs.184052 PP1201 protein /FL=gb:NM_022152.1	NM_022152		NP_071435

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217731_s_at	0.041055	gb:NM_021999.1 /DEF=Homo sapiens integral membrane protein 2B (ITM2B), mRNA. /FEA=mRNA /GEN=ITM2B /PROD=integral membrane protein 2B /DB_XREF=gi:11527401 /UG=Hs.239625 integral membrane protein 2B /FL=gb:NM_021999.1 gb:AF136973.1 gb:BC000554.1 gb:AF092128.1 gb:AF152462.1 gb:AF246221.1	NM_021999		NP_068839
217736_s_at	5.03E-04	gb:NM_014413.2 /DEF=Homo sapiens heme-regulated initiation factor 2-alpha kinase (HRI), mRNA. /FEA=mRNA /GEN=HRI /PROD=heme-regulated initiation factor 2-alpha kinase /DB_XREF=gi:11125767 /UG=Hs.258730 heme-regulated initiation factor 2-alpha kinase /FL=gb:NM_014413.2 gb:AL136563.1 gb:AF147094.1 gb:AF255050.1 gb:AF116634.1 gb:AF183414.1	NM_014413		NP_055228
217737_x_at	1.26E-04	gb:NM_016407.1 /DEF=Homo sapiens hypothetical protein (HSPC164), mRNA. /FEA=mRNA /GEN=HSPC164 /PROD=hypothetical protein /DB_XREF=gi:7705482 /UG=Hs.182281 hypothetical protein /FL=gb:BC003359.1 gb:AF212244.1 gb:AF117231.1 gb:AF161513.1 gb:AF161518.1 gb:NM_016407.1	NM_016407		NP_057491
217739_s_at	0.033381	gb:NM_005746.1 /DEF=Homo sapiens pre-B-cell colony-enhancing factor (PBEF), mRNA. /FEA=mRNA /GEN=PBEF /PROD=pre-B-cell colony-enhancing factor /DB_XREF=gi:5031976 /UG=Hs.239138 pre-B-cell colony-enhancing factor /FL=gb:U02020.1 gb:NM_005746.1	NM_005746		NP_005737
217741_s_at	3.26E-04	zinc finger protein 216	AW471220	Hs.3776	NP_005998

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217742_s_at	0.004341	gb:NM_016628.1 /DEF=Homo sapiens hypothetical protein (LOC51322), mRNA. /FEA=mRNA /GEN=LOC51322 /PROD=hypothetical protein /DB_XREF=gi:7706169 /UG=Hs.70333 hypothetical protein /FL=gb:BC004258.1 gb:AF208858.1 gb:NM_016628.1	NM_016628		NP_567823
217743_s_at	1.26E-04	gb:NM_018247.1 /DEF=Homo sapiens hypothetical protein FLJ10856 (FLJ10856), mRNA. /FEA=mRNA /GEN=FLJ10856 /PROD=hypothetical protein FLJ10856 /DB_XREF=gi:8922719 /UG=Hs.108530 hypothetical protein FLJ10856 /FL=gb:NM_018247.1	NM_018247		NP_060717
217745_s_at	0.001116	gb:NM_025146.1 /DEF=Homo sapiens hypothetical protein FLJ13194 (FLJ13194), mRNA. /FEA=mRNA /GEN=FLJ13194 /PROD=hypothetical protein FLJ13194 /DB_XREF=gi:13376734 /UG=Hs.288932 hypothetical protein FLJ13194 /FL=gb:NM_025146.1	NM_025146		NP_079422
217746_s_at	1.26E-04	gb:NM_013374.1 /DEF=Homo sapiens programmed cell death 6 interacting protein (PDCD6IP), mRNA. /FEA=mRNA /GEN=PDCD6IP /PROD=programmed cell death 6-interacting protein /DB_XREF=gi:7019486 /UG=Hs.9663 programmed cell death 6-interacting protein /FL=gb:AF349951.1 gb:AF151793.1 gb:NM_013374.1	NM_013374		NP_037506
217747_s_at	0.041055	gb:NM_001013.1 /DEF=Homo sapiens ribosomal protein S9 (RPS9), mRNA. /FEA=mRNA /GEN=RPS9 /PROD=ribosomal protein S9 /DB_XREF=gi:4506744 /UG=Hs.180920 ribosomal protein S9 /FL=gb:BC000802.1 gb:NM_001013.1 gb:U14971.1	NM_001013		NP_001004

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217748_at	1.26E-04	gb:NM_015999.1 /DEF=Homo sapiens CGI-45 protein (LOC51094), mRNA. /FEA=mRNA /GEN=LOC51094 /PROD=CGI-45 protein /DB_XREF=gi:7705760 /UG=Hs.5298 CGI-45 protein /FL=gb:AF151803.1 gb:NM_015999.1	NM_015999		NP_057083
217755_at	1.26E-04	gb:NM_016185.1 /DEF=Homo sapiens hematological and neurological expressed 1 (HN1), mRNA. /FEA=mRNA /GEN=HN1 /PROD=hematological and neurological expressed 1 /DB_XREF=gi:7705876 /UG=Hs.109706 hematological and neurological expressed 1 /FL=gb:AF060925.1 gb:BC001420.1 gb:AF177862.1 gb:NM_016185.1	NM_016185		NP_057269
217758_s_at	0.00161	gb:NM_020123.1 /DEF=Homo sapiens endomembrane protein emp70 precursor isolog (LOC56889), mRNA. /FEA=mRNA /GEN=LOC56889 /PROD=endomembrane protein emp70 precursor isolog /DB_XREF=gi:10047129 /UG=Hs.8203 endomembrane protein emp70 precursor isolog /FL=gb:NM_020123.1 gb:AF160213.1 gb:AF269150.1	NM_020123		NP_064508
217762_s_at	1.26E-04	RAB31, member RAS oncogene family	BE789881	Hs.223025	NP_006859
217764_s_at	1.26E-04	gb:AF183421.1 /DEF=Homo sapiens small GTP-binding protein rab22b mRNA, complete cds. /FEA=mRNA /PROD=small GTP-binding protein rab22b /DB_XREF=gi:9963780 /UG=Hs.223025 RAB31, member RAS oncogene family /FL=gb:AF234995.1 gb:BC001148.1 gb:U59877.1 gb:U57091.1 gb:NM_006868.1 gb:AF183421.1	AF183421		NP_006859

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217766_s_at	0.033381	gb:NM_014313.1 /DEF=Homo sapiens small membrane protein 1 (SMP1), mRNA. /FEA=mRNA /GEN=SMP1 /PROD=small membrane protein 1 /DB_XREF=gi:7657594 /UG=Hs.107979 small membrane protein 1 /FL=gb:AL136627.1 gb:AF081282.1 gb:NM_014313.1	NM_014313		NP_055128
217768_at	0.026891	gb:NM_016039.1 /DEF=Homo sapiens CGI-99 protein (LOC51637), mRNA. /FEA=mRNA /GEN=LOC51637 /PROD=CGI-99 protein /DB_XREF=gi:7706321 /UG=Hs.110803 CGI-99 protein /FL=gb:BC001722.1 gb:AF151857.1 gb:AF100755.1 gb:NM_016039.1	NM_016039		NP_057123
217769_s_at	0.004341	gb:NM_015932.1 /DEF=Homo sapiens hypothetical protein (HSPC014), mRNA. /FEA=mRNA /GEN=HSPC014 /PROD=hypothetical protein /DB_XREF=gi:7705428 /UG=Hs.279813 hypothetical protein /FL=gb:AF275807.1 gb:BC003390.1 gb:AF077200.1 gb:AF125097.1 gb:NM_015932.1	NM_015932		NP_057016
217772_s_at	0.004733	gb:NM_014342.1 /DEF=Homo sapiens mitochondrial carrier homolog 2 (MTCH2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MTCH2 /PROD=mitochondrial carrier homolog 2 /DB_XREF=gi:7657346 /UG=Hs.279609 mitochondrial carrier homolog 2 /FL=gb:BC000875.1 gb:AF085361.1 gb:AF176008.1 gb:NM_014342.1	NM_014342		NP_055157

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217774_s_at	0.026891	gb:NM_016404.1 /DEF=Homo sapiens hypothetical protein (HSPC152), mRNA. /FEA=mRNA /GEN=HSPC152 /PROD=hypothetical protein /DB_XREF=gi:7705476 /UG=Hs.79259 hypothetical protein /FL=gb:AF110774.1 gb:AF161501.1 gb:NM_016404.1 gb:AF229068.1	NM_016404		NP_057488
217780_at	0.041055	gb:NM_016145.1 /DEF=Homo sapiens PTD008 protein (PTD008), mRNA. /FEA=mRNA /GEN=PTD008 /PROD=PTD008 protein /DB_XREF=gi:7706664 /UG=Hs.108969 PTD008 protein /FL=gb:AF059620.1 gb:AF151898.1 gb:AF078861.1 gb:NM_016145.1	NM_016145		NP_057229
217781_s_at	1.26E-04	gb:NM_022473.1 /DEF=Homo sapiens zinc finger protein 106 (ZFP106), mRNA. /FEA=mRNA /GEN=ZFP106 /PROD=zinc finger protein 106 /DB_XREF=gi:11968022 /UG=Hs.15220 zinc finger protein 106 /FL=gb:AF205632.1 gb:NM_022473.1	NM_022473		NP_071918
217783_s_at	0.013223	gb:NM_016061.1 /DEF=Homo sapiens CGI-127 protein (LOC51646), mRNA. /FEA=mRNA /GEN=LOC51646 /PROD=CGI-127 protein /DB_XREF=gi:7706340 /UG=Hs.184542 CGI-127 protein /FL=gb:BC000836.1 gb:AF151885.1 gb:NM_016061.1	NM_016061		NP_057145
217786_at	9.79E-04	gb:NM_006109.1 /DEF=Homo sapiens skb1 (S. pombe) homolog (SKB1), mRNA. /FEA=mRNA /GEN=SKB1 /PROD=skb1 (S. pombe) homolog /DB_XREF=gi:5174682 /UG=Hs.12912 skb1 (S. pombe) homolog /FL=gb:AF015913.1 gb:NM_006109.1 gb:AF167572.1	NM_006109		NP_006100

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217788_s_at	7.58E-04	gb:NM_004481.2 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2) (GALNT2), mRNA. /FEA=mRNA /GEN=GALNT2 /PROD=polypeptide N-acetylgalactosaminyltransferase 2 /DB_XREF=gi:9945385 /UG=Hs.130181 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2) /FL=gb:NM_004481.2	NM_004481		NP_004472
217794_at	7.58E-04	gb:NM_018457.1 /DEF=Homo sapiens DKFZp564J157 protein (DKFZP564J157), mRNA. /FEA=mRNA /GEN=DKFZP564J157 /PROD=DKFZp564J157 protein /DB_XREF=gi:8922156 /UG=Hs.63042 DKFZp564J157 protein /FL=gb:AF217517.1 gb:NM_018457.1	NM_018457		NP_060927
217795_s_at	0.005852	hypothetical protein MGC3222	W74580	Hs.323193	NP_077310
217797_at	0.007779	gb:NM_016406.1 /DEF=Homo sapiens hypothetical protein (HSPC155), mRNA. /FEA=mRNA /GEN=HSPC155 /PROD=hypothetical protein /DB_XREF=gi:7705480 /UG=Hs.177507 hypothetical protein /FL=gb:BC005187.1 gb:AF151884.1 gb:AF161504.1 gb:NM_016406.1	NM_016406		NP_057490
217799_x_at	0.027239	gb:NM_003344.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2H (homologous to yeast UBC8) (UBE2H), mRNA. /FEA=mRNA /GEN=UBE2H /PROD=ubiquitin-conjugating enzyme E2H (homologous to yeast UBC8) /DB_XREF=gi:4507782 /UG=Hs.28505 ubiquitin-conjugating enzyme E2H (homologous to yeast UBC8) /FL=gb:NM_003344.1	NM_003344		NP_003335

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217800_s_at	0.001116	gb:NM_030571.1 /DEF=Homo sapiens hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5 (MGC10924), mRNA. /FEA=mRNA /GEN=MGC10924 /PROD=hypothetical protein MGC10924 similar to Nedd4WW-binding protein 5 /DB_XREF=gi:13386479 /UG=Hs.9788 hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5 /FL=gb:BC004317.1 gb:NM_030571.1	NM_030571		NP_085048
217801_at	0.003171	gb:NM_006886.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1 complex, epsilon subunit (ATP5E), mRNA. /FEA=mRNA /GEN=ATP5E /PROD=ATP synthase, H ⁺ transporting, mitochondrial F1complex, epsilon subunit /DB_XREF=gi:5901895 /UG=Hs.177530 ATP synthase, H ⁺ transporting, mitochondrial F1 complex, epsilon subunit /FL=gb:BC001690.1 gb:BC003671.1 gb:AF077045.1 gb:NM_006886.1 gb:AF052955.1	NM_006886		NP_008817
217802_s_at	1.26E-04	gb:NM_022731.1 /DEF=Homo sapiens similar to rat nuclear ubiquitous casein kinase 2 (NUCKS), mRNA. /FEA=mRNA /GEN=NUCKS /PROD=similar to rat nuclear ubiquitous casein kinase2 /DB_XREF=gi:12232386 /UG=Hs.118064 similar to rat nuclear ubiquitous casein kinase 2 /FL=gb:NM_022731.1 gb:BC000805.1	NM_022731		NP_073568
217803_at	1.26E-04	gb:NM_022130.1 /DEF=Homo sapiens Golgi protein (GPP34), mRNA. /FEA=mRNA /GEN=GPP34 /PROD=Golgi protein /DB_XREF=gi:11545858 /UG=Hs.18271 Golgi protein /FL=gb:NM_022130.1	NM_022130		NP_071413

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217805_at	0.003171	gb:NM_004516.1 /DEF=Homo sapiens interleukin enhancer binding factor 3, 90kD (ILF3), mRNA. /FEA=mRNA /GEN=ILF3 /PROD=interleukin enhancer binding factor 3, 90kD /DB_XREF=gi:4758603 /UG=Hs.256583 interleukin enhancer binding factor 3, 90kD /FL=gb:BC003086.1 gb:NM_004516.1 gb:U10324.1 gb:AF167570.1 gb:NM_012218.1	NM_004516		NP_703194
217807_s_at	0.016934	gb:NM_015710.1 /DEF=Homo sapiens glioma tumor suppressor candidate region gene 2 (GLTSCR2), mRNA. /FEA=mRNA /GEN=GLTSCR2 /PROD=glioma tumor suppressor candidate region gene 2 /DB_XREF=gi:7657129 /UG=Hs.2237 glioma tumor suppressor candidate region gene 2. /FL=gb:AF182076.1 gb:NM_015710.1	NM_015710		NP_056525
217811_at	1.26E-04	gb:NM_016275.1 /DEF=Homo sapiens selenoprotein T (LOC51714), mRNA. /FEA=mRNA /GEN=LOC51714 /PROD=selenoprotein T /DB_XREF=gi:7706470 /UG=Hs.8148 selenoprotein T /FL=gb:AF131856.1 gb:AF195141.1 gb:NM_016275.1	NM_016275		NP_057359
217812_at	0.016934	gb:NM_016258.1 /DEF=Homo sapiens high-glucose-regulated protein 8 (HGRG8), mRNA. /FEA=mRNA /GEN=HGRG8 /PROD=high-glucose-regulated protein 8 /DB_XREF=gi:7705410 /UG=Hs.20993 high-glucose-regulated protein 8 /FL=gb:BC002559.1 gb:AF155095.1 gb:AF192968.1 gb:NM_016258.1	NM_016258		NP_057342

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217814_at	1.26E-04	gb:NM_020198.1 /DEF=Homo sapiens GK001 protein (GK001), mRNA. /FEA=mRNA /GEN=GK001 /PROD=GK001 protein /DB_XREF=gi:9910241 /UG=Hs.8207 GK001 protein /FL=gb:AF113221.1 gb:BC001300.1 gb:AF226054.1 gb:NM_020198.1	NM_020198		NP_064583
217815_at	0.00161	gb:NM_007192.1 /DEF=Homo sapiens chromatin-specific transcription elongation factor, 140 kDa subunit (FACTP140), mRNA. /FEA=mRNA /GEN=FACTP140 /PROD=chromatin-specific transcription elongationfactor, 140 kDa subunit /DB_XREF=gi:6005756 /UG=Hs.14963 chromatin-specific transcription elongation factor, 140 kDa subunit /FL=gb:AF152961.1 gb:NM_007192.1	NM_007192		NP_009123
217819_at	1.26E-04	gb:NM_016099.1 /DEF=Homo sapiens HSPC041 protein (LOC51125), mRNA. /FEA=mRNA /GEN=LOC51125 /PROD=HSPC041 protein /DB_XREF=gi:7705820 /UG=Hs.7953 HSPC041 protein /FL=gb:AF125102.1 gb:NM_016099.1	NM_016099		NP_057183
217823_s_at	1.26E-04	Consensus includes gb:AL562528 /FEA=EST /DB_XREF=gi:12911037 /DB_XREF=est:AL562528 /CLONE=CS0DC012YI08 (3 prime) /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	NM_016021		NP_057420

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217824_at	0.02145	Consensus includes gb:AW500009 /FEA=EST /DB_XREF=gi:7112213 /DB_XREF=est:UI-HF-BN0-aki-e-09-0-UI.r1 /CLONE=IMAGE:3077105 /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	NM_016021		NP_057420
217825_s_at	0.00228	gb:AF151039.1 /DEF=Homo sapiens HSPC205 mRNA, complete cds. /FEA=mRNA /PROD=HSPC205 /DB_XREF=gi:7106799 /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	AF151039		NP_057420
217826_s_at	7.58E-04	gb:NM_016021.1 /DEF=Homo sapiens CGI-76 protein (LOC51632), mRNA. /FEA=mRNA /GEN=LOC51632 /PROD=CGI-76 protein /DB_XREF=gi:7706311 /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	NM_016021		NP_057420
217827_s_at	2.05E-04	gb:NM_016630.2 /DEF=Homo sapiens acid cluster protein 33 (ACP33), mRNA. /FEA=mRNA /GEN=ACP33 /PROD=acid cluster protein 33 /DB_XREF=gi:13699804 /UG=Hs.71475 acid cluster protein 33 /FL=gb:BC000244.1 gb:NM_016630.2 gb:AF208861.1	NM_016630		NP_057714
217835_x_at	3.26E-04	gb:NM_018840.1 /DEF=Homo sapiens putative Rab5-interacting protein (LOC55969), mRNA. /FEA=mRNA /GEN=LOC55969 /PROD=putative Rab5-interacting protein /DB_XREF=gi:10047115 /UG=Hs.184062 putative Rab5-interacting protein /FL=gb:NM_018840.1 gb:AF274936.1 gb:AF112213.1	NM_018840		NP_061328

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217836_s_at	0.005852	gb:NM_018253.1 /DEF=Homo sapiens hypothetical protein FLJ10875 (FLJ10875), mRNA. /FEA=mRNA /GEN=FLJ10875 /PROD=hypothetical protein FLJ10875 /DB_XREF=gi:8922730 /UG=Hs.18851 hypothetical protein FLJ10875 /FL=gb:BC001655.1 gb:BC001843.1 gb:NM_018253.1	NM_018253		NP_620832
217837_s_at	1.26E-04	gb:NM_016079.1 /DEF=Homo sapiens CGI-149 protein (LOC51652), mRNA. /FEA=mRNA /GEN=LOC51652 /PROD=CGI-149 protein /DB_XREF=gi:7706352 /UG=Hs.189658 CGI-149 protein /FL=gb:BC004419.1 gb:AF151907.1 gb:AF219226.1 gb:NM_016079.1	NM_016079		NP_057163
217838_s_at	0.033381	gb:NM_016337.1 /DEF=Homo sapiens RNB6 (RNB6), mRNA. /FEA=mRNA /GEN=RNB6 /PROD=RNB6 /DB_XREF=gi:7706686 /UG=Hs.241471 RNB6 /FL=gb:AF052504.1 gb:NM_016337.1	NM_016337		NP_057421
217839_at	6.42E-04	gb:NM_006070.1 /DEF=Homo sapiens TRK-fused gene (TFG), mRNA. /FEA=mRNA /GEN=TFG /PROD=TRK-fused gene /DB_XREF=gi:5174718 /UG=Hs.250897 TRK-fused gene /FL=gb:NM_006070.1	NM_006070		NP_006061
217843_s_at	3.26E-04	gb:NM_014166.1 /DEF=Homo sapiens HSPC126 protein (HSPC126), mRNA. /FEA=mRNA /GEN=HSPC126 /PROD=HSPC126 protein /DB_XREF=gi:7661787 /UG=Hs.181112 HSPC126 protein /FL=gb:AF300618.1 gb:BC005189.1 gb:AF161475.1 gb:AF230381.1 gb:NM_014166.1	NM_014166		NP_054885

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217844_at	0.003529	gb:NM_021198.1 /DEF=Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA. /FEA=mRNA /GEN=NLI-IF /PROD=nuclear LIM interactor-interacting factor /DB_XREF=gi:10864008 /UG=Hs.283724 nuclear LIM interactor-interacting factor /FL=gb:AF229162.1 gb:NM_021198.1	NM_021198		NP_067021
217845_x_at	0.00161	gb:NM_014056.1 /DEF=Homo sapiens DKFZP564K247 protein (DKFZP564K247), mRNA. /FEA=mRNA /GEN=DKFZP564K247 /PROD=DKFZP564K247 protein /DB_XREF=gi:7661619 /UG=Hs.7917 DKFZP564K247 protein /FL=gb:BC000601.1 gb:AF077034.1 gb:AF145385.1 gb:AL110233.1 gb:NM_014056.1	NM_014056		NP_054775
217852_s_at	1.26E-04	gb:NM_018184.1 /DEF=Homo sapiens hypothetical protein FLJ10702 (FLJ10702), mRNA. /FEA=mRNA /GEN=FLJ10702 /PROD=hypothetical protein FLJ10702 /DB_XREF=gi:8922600 /UG=Hs.104222 hypothetical protein FLJ10702 /FL=gb:NM_018184.1	NM_018184		NP_060654
217854_s_at	0.022795	gb:NM_002695.1 /DEF=Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E (25kD) (POLR2E), mRNA. /FEA=mRNA /GEN=POLR2E /PROD=polymerase (RNA) II (DNA directed) polypeptide E(25kD) /DB_XREF=gi:4505944 /UG=Hs.24301 polymerase (RNA) II (DNA directed) polypeptide E (25kD) /FL=gb:BC004441.1 gb:NM_002695.1 gb:J04965.1 gb:D38251.1	NM_002695		NP_002686

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217855_x_at	0.004341	gb:NM_016547.1 /DEF=Homo sapiens calcium binding protein Cab45 precursor, (LOC51767), mRNA. /FEA=mRNA /GEN=LOC51767 /PROD=calcium binding protein Cab45 precursor, /DB_XREF=gi:7706572 /UG=Hs.42806 calcium binding protein Cab45 precursor, /FL=gb:AF178986.1 gb:NM_016547.1	NM_016547		NP_057631
217862_at	3.26E-04	Consensus includes gb:N24868 /FEA=EST /DB_XREF=gi:1139018 /DB_XREF=est:yx70h06.s1 /CLONE=IMAGE:267131 /UG=Hs.75251 DEADH (Asp-Glu-Ala-AspHis) box binding protein 1 /FL=gb:AF077951.1 gb:AF167160.1 gb:NM_016166.1	NM_016166		NP_057250
217863_at	0.00228	Consensus includes gb:AI348378 /FEA=EST /DB_XREF=gi:4085584 /DB_XREF=est:qo20h06.x1 /CLONE=IMAGE:1909115 /UG=Hs.75251 DEADH (Asp-Glu-Ala-AspHis) box binding protein 1 /FL=gb:AF077951.1 gb:AF167160.1 gb:NM_016166.1	NM_016166		NP_057250
217865_at	2.05E-04	gb:NM_018434.1 /DEF=Homo sapiens goliath protein (LOC55819), mRNA. /FEA=mRNA /GEN=LOC55819 /PROD=goliath protein /DB_XREF=gi:10092650 /UG=Hs.102737 goliath protein /FL=gb:NM_018434.1	NM_018434		NP_060904
217868_s_at	1.26E-04	gb:NM_016025.1 /DEF=Homo sapiens CGI-81 protein (DREV1), mRNA. /FEA=mRNA /GEN=DREV1 /PROD=CGI-81 protein /DB_XREF=gi:7705788 /UG=Hs.279583 CGI-81 protein /FL=gb:BC000195.1 gb:AF151839.1 gb:NM_016025.1	NM_016025		NP_057109

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217869_at	1.26E-04	gb:NM_016142.1 /DEF=Homo sapiens steroid dehydrogenase homolog (LOC51144), mRNA. /FEA=mRNA /GEN=LOC51144 /PROD=steroid dehydrogenase homolog /DB_XREF=gi:7705854 /UG=Hs.279617 steroid dehydrogenase homolog /FL=gb:AF078850.1 gb:NM_016142.1	NM_016142		NP_057226
217873_at	1.26E-04	gb:NM_016289.1 /DEF=Homo sapiens MO25 protein (LOC51719), mRNA. /FEA=mRNA /GEN=LOC51719 /PROD=MO25 protein /DB_XREF=gi:7706480 /UG=Hs.6406 MO25 protein /FL=gb:AF113536.1 gb:NM_016289.1	NM_016289		NP_057373
217874_at	1.26E-04	gb:NM_003849.1 /DEF=Homo sapiens succinate-CoA ligase, GDP-forming, alpha subunit (SUCLG1), mRNA. /FEA=mRNA /GEN=SUCLG1 /PROD=succinate-CoA ligase, GDP-forming, alphasubunit /DB_XREF=gi:11321580 /UG=Hs.7043 succinate-CoA ligase, GDP-forming, alpha subunit /FL=gb:NM_003849.1 gb:BC000504.1 gb:AF104921.2	NM_003849		NP_003840
217877_s_at	0.02145	gb:NM_021639.1 /DEF=Homo sapiens hypothetical protein SP192 (SP192), mRNA. /FEA=mRNA /GEN=SP192 /PROD=hypothetical protein SP192 /DB_XREF=gi:11056015 /UG=Hs.169854 hypothetical protein SP192 /FL=gb:NM_021639.1	NM_021639		NP_067652
217880_at	0.005852	cell division cycle 27	AI203880	Hs.408978	NP_001247
217882_at	0.00161	gb:NM_018447.1 /DEF=Homo sapiens 30 kDa protein (LOC55831), mRNA. /FEA=mRNA /GEN=LOC55831 /PROD=30 kDa protein /DB_XREF=gi:8923856 /UG=Hs.283714 30 kDa protein /FL=gb:AF157321.1 gb:NM_018447.1	NM_018447		NP_060917

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217883_at	1.26E-04	gb:NM_015702.1 /DEF=Homo sapiens hypothetical protein (CL25022), mRNA. /FEA=mRNA /GEN=CL25022 /PROD=hypothetical protein /DB_XREF=gi:7661547 /UG=Hs.5324 hypothetical protein /FL=gb:AF060224.1 gb:BC000932.2 gb:AF131802.1 gb:AF161510.1 gb:NM_015702.1	NM_015702		NP_056517
217885_at	3.26E-04	gb:NM_018085.1 /DEF=Homo sapiens hypothetical protein FLJ10402 (FLJ10402), mRNA. /FEA=mRNA /GEN=FLJ10402 /PROD=hypothetical protein FLJ10402 /DB_XREF=gi:8922404 /UG=Hs.7579 hypothetical protein FLJ10402 /FL=gb:BC003604.1 gb:NM_018085.1	NM_018085		NP_060555
217886_at	0.001116	epidermal growth factor receptor pathway substrate 15	BF213575	Hs.389343	NP_001972
217887_s_at	0.007779	gb:NM_001981.1 /DEF=Homo sapiens epidermal growth factor receptor pathway substrate 15 (EPS15), mRNA. /FEA=mRNA /GEN=EPS15 /PROD=epidermal growth factor receptor pathway substrate 15 /DB_XREF=gi:4503592 /UG=Hs.79095 epidermal growth factor receptor pathway substrate 15 /FL=gb:NM_001981.1 gb:U07707.1	NM_001981		NP_001972
217898_at	2.05E-04	gb:NM_020154.1 /DEF=Homo sapiens chromosome 11 hypothetical protein ORF3 (LOC56851), mRNA. /FEA=mRNA /GEN=LOC56851 /PROD=chromosome 11 hypothetical protein ORF3 /DB_XREF=gi:9910345 /UG=Hs.4245 chromosome 11 hypothetical protein ORF3 /FL=gb:AF242729.1 gb:NM_020154.1	NM_020154		NP_064539

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217900_at	0.005852	gb:NM_018060.1 /DEF=Homo sapiens hypothetical protein FLJ10326 (FLJ10326), mRNA. /FEA=mRNA /GEN=FLJ10326 /PROD=hypothetical protein FLJ10326 /DB_XREF=gi:8922355 /UG=Hs.262823 hypothetical protein FLJ10326 /FL=gb:NM_018060.1	NM_018060		NP_060530
217903_at	0.016934	gb:NM_013403.1 /DEF=Homo sapiens zinedin (ZIN), mRNA. /FEA=mRNA /GEN=ZIN /PROD=zinedin /DB_XREF=gi:7019572 /UG=Hs.108665 zinedin /FL=gb:AF212940.1 gb:NM_013403.1	NM_013403		NP_037535
217905_at	7.58E-04	gb:NM_024834.1 /DEF=Homo sapiens hypothetical protein FLJ13081 (FLJ13081), mRNA. /FEA=mRNA /GEN=FLJ13081 /PROD=hypothetical protein FLJ13081 /DB_XREF=gi:13376242 /UG=Hs.180638 hypothetical protein FLJ13081 /FL=gb:BC004183.1 gb:NM_024834.1	NM_024834		NP_079110
217907_at	7.58E-04	gb:NM_014161.1 /DEF=Homo sapiens HSPC071 protein (HSPC071), mRNA. /FEA=mRNA /GEN=HSPC071 /PROD=HSPC071 protein /DB_XREF=gi:7661777 /UG=Hs.23038 HSPC071 protein /FL=gb:AL136633.1 gb:BC001623.1 gb:AF161556.1 gb:NM_014161.1	NM_014161		NP_054880
217908_s_at	1.26E-04	gb:NM_018442.1 /DEF=Homo sapiens PC326 protein (PC326), mRNA. /FEA=mRNA /GEN=PC326 /PROD=PC326 protein /DB_XREF=gi:8923955 /UG=Hs.279882 PC326 protein /FL=gb:AL136738.1 gb:AF150734.1 gb:NM_018442.1	NM_018442		NP_060912

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217910_x_at	1.26E-04	gb:NM_013383.1 /DEF=Homo sapiens MAX-like bHLHZIP protein (MLX), mRNA. /FEA=mRNA /GEN=MLX /PROD=MAX-like bHLHZIP protein /DB_XREF=gi:7019460 /UG=Hs.78185 MAX-like bHLHZIP protein /FL=gb:AF213668.1 gb:AF203978.1 gb:NM_013383.1	NM_013383		NP_733753
217911_s_at	0.003171	gb:NM_004281.1 /DEF=Homo sapiens BCL2-associated athanogene 3 (BAG3), mRNA. /FEA=mRNA /GEN=BAG3 /PROD=BCL2-associated athanogene 3 /DB_XREF=gi:6631072 /UG=Hs.15259 BCL2-associated athanogene 3 /FL=gb:AF095193.2 gb:NM_004281.1 gb:AF127139.1 gb:AF071218.2	NM_004281		NP_004272
217913_at	0.013223	gb:NM_013245.1 /DEF=Homo sapiens vacuolar sorting protein 4 (VPS4), mRNA. /FEA=mRNA /GEN=VPS4 /PROD=vacuolar sorting protein 4 /DB_XREF=gi:7019568 /UG=Hs.234839 vacuolar sorting protein 4 /FL=gb:AF159063.1 gb:AF112215.1 gb:NM_013245.1	NM_013245		NP_037377
217915_s_at	0.016934	gb:NM_016304.1 /DEF=Homo sapiens 60S ribosomal protein L30 isolog (LOC51187), mRNA. /FEA=mRNA /GEN=LOC51187 /PROD=60S ribosomal protein L30 isolog /DB_XREF=gi:10047101 /UG=Hs.284162 60S ribosomal protein L30 isolog /FL=gb:NM_016304.1 gb:AF060926.1 gb:AF212226.1 gb:BC005344.1 gb:AF201949.1 gb:AF165521.1	NM_016304		NP_057388

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217916_s_at	0.00161	gb:NM_016623.1 /DEF=Homo sapiens hypothetical protein (BM-009), mRNA. /FEA=mRNA /GEN=BM-009 /PROD=hypothetical protein /DB_XREF=gi:7705303 /UG=Hs.92918 hypothetical protein /FL=gb:BC003599.1 gb:AF208851.1 gb:NM_016623.1	NM_016623		NP_057707
217917_s_at	5.03E-04	gb:NM_014183.1 /DEF=Homo sapiens HSPC162 protein (HSPC162), mRNA. /FEA=mRNA /GEN=HSPC162 /PROD=HSPC162 protein /DB_XREF=gi:7661821 /UG=Hs.100002 HSPC162 protein /FL=gb:BC002481.1 gb:AY026513.1 gb:AF161511.1 gb:NM_014183.1 gb:AF165516.1	NM_014183		NP_808853
217918_at	1.26E-04	gb:NM_014183.1 /DEF=Homo sapiens HSPC162 protein (HSPC162), mRNA. /FEA=mRNA /GEN=HSPC162 /PROD=HSPC162 protein /DB_XREF=gi:7661821 /UG=Hs.100002 HSPC162 protein /FL=gb:BC002481.1 gb:AY026513.1 gb:AF161511.1 gb:NM_014183.1 gb:AF165516.1	NM_014183		NP_808853
217923_at	0.02145	gb:NM_012392.1 /DEF=Homo sapiens PEF protein with a long N-terminal hydrophobic domain (peflin) (PEF), mRNA. /FEA=mRNA /GEN=PEF /PROD=peflin /DB_XREF=gi:6912581 /UG=Hs.241531 peflin /FL=gb:BC002773.1 gb:AB018357.1 gb:AB026628.1 gb:NM_012392.1	NM_012392		NP_036524
217925_s_at	0.022752	gb:NM_022758.1 /DEF=Homo sapiens hypothetical protein FLJ22195 (FLJ22195), mRNA. /FEA=mRNA /GEN=FLJ22195 /PROD=hypothetical protein FLJ22195 /DB_XREF=gi:12232426 /UG=Hs.25999 hypothetical protein FLJ22195 /FL=gb:NM_022758.1	NM_022758		NP_073595

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217927_at	3.26E-04	gb:NM_014041.1 /DEF=Homo sapiens HSPC033 protein (HSPC033), mRNA. /FEA=mRNA /GEN=HSPC033 /PROD=HSPC033 protein /DB_XREF=gi:7661745 /UG=Hs.11125 HSPC033 protein /FL=gb:BC000884.1 gb:AF092138.1 gb:NM_014041.1	NM_014041		NP_054760
217928_s_at	0.041055	gb:NM_018312.2 /DEF=Homo sapiens chromosome 11 open reading frame 23 (C11ORF23), mRNA. /FEA=mRNA /GEN=C11ORF23 /PROD=sporulation-induced transcript 4-associated protein /DB_XREF=gi:13489082 /UG=Hs.180817 chromosome 11 open reading frame 23 /FL=gb:AF264779.1 gb:NM_018312.2	NM_018312		NP_060782
217930_s_at	0.009689	gb:NM_019009.1 /DEF=Homo sapiens TOLLIP protein (LOC54472), mRNA. /FEA=mRNA /GEN=LOC54472 /PROD=TOLLIP protein /DB_XREF=gi:9506854 /UG=Hs.25413 TOLLIP protein /FL=gb:AL136835.1 gb:BC004420.1 gb:NM_019009.1	NM_019009		NP_061882
217936_at	0.003171	Rho GTPase activating protein 5	AW044631	Hs.267831	
217938_s_at	0.00228	gb:NM_020122.1 /DEF=Homo sapiens potassium channel modulatory factor (DKFZP434L1021), mRNA. /FEA=mRNA /GEN=DKFZP434L1021 /PROD=potassium channel modulatory factor /DB_XREF=gi:10047127 /UG=Hs.5392 potassium channel modulatory factor /FL=gb:NM_020122.1 gb:BC000178.2	NM_020122		NP_064507

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217939_s_at	7.58E-04	gb:NM_017657.1 /DEF=Homo sapiens hypothetical protein FLJ20080 (FLJ20080), mRNA. /FEA=mRNA /GEN=FLJ20080 /PROD=hypothetical protein FLJ20080 /DB_XREF=gi:8923087 /UG=Hs.7942 hypothetical protein FLJ20080 /FL=gb:NM_017657.1	NM_017657		NP_060127
217941_s_at	0.00228	gb:NM_018695.1 /DEF=Homo sapiens erbb2-interacting protein ERBIN (LOC55914), mRNA. /FEA=mRNA /GEN=LOC55914 /PROD=erbb2-interacting protein ERBIN /DB_XREF=gi:8923908 /UG=Hs.8117 erbb2-interacting protein ERBIN /FL=gb:AF263744.1 gb:NM_018695.1	NM_018695		NP_061165
217949_s_at	5.03E-04	gb:NM_024006.1 /DEF=Homo sapiens hypothetical protein IMAGE3455200 (IMAGE3455200), mRNA. /FEA=mRNA /GEN=IMAGE3455200 /PROD=hypothetical protein IMAGE3455200 /DB_XREF=gi:13124769 /UG=Hs.324844 hypothetical protein IMAGE3455200 /FL=gb:BC002911.1 gb:NM_024006.1	NM_024006		NP_076869
217951_s_at	5.03E-04	PHD finger protein 3	AW189430	Hs.78893	NP_055968
217952_x_at	0.016934	PHD finger protein 3	AW189430	Hs.78893	NP_055968
217954_s_at	0.026891	gb:NM_015153.1 /DEF=Homo sapiens KIAA0244 protein (KIAA0244), mRNA. /FEA=mRNA /GEN=KIAA0244 /PROD=KIAA0244 protein /DB_XREF=gi:7662017 /UG=Hs.78893 KIAA0244 protein /FL=gb:AF091622.1 gb:NM_015153.1	NM_015153		NP_055968

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217955_at	1.26E-04	gb:NM_015367.1 /DEF=Homo sapiens MIL1 protein (MIL1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MIL1 /PROD=MIL1 protein /DB_XREF=gi:7662505 /UG=Hs.10267 MIL1 protein /FL=gb:AF325209.1 gb:AF146568.1 gb:NM_015367.1	NM_015367		NP_056182
217958_at	0.026891	gb:NM_016146.1 /DEF=Homo sapiens PTD009 protein (PTD009), mRNA. /FEA=mRNA /GEN=PTD009 /PROD=PTD009 protein /DB_XREF=gi:7706666 /UG=Hs.279901 PTD009 protein /FL=gb:AF151862.1 gb:AF078862.1 gb:AF161520.1 gb:NM_016146.1	NM_016146		NP_057230
217959_s_at	0.004341	gb:NM_016146.1 /DEF=Homo sapiens PTD009 protein (PTD009), mRNA. /FEA=mRNA /GEN=PTD009 /PROD=PTD009 protein /DB_XREF=gi:7706666 /UG=Hs.279901 PTD009 protein /FL=gb:AF151862.1 gb:AF078862.1 gb:AF161520.1 gb:NM_016146.1	NM_016146		NP_057230
217962_at	1.26E-04	gb:NM_018648.1 /DEF=Homo sapiens nucleolar protein family A, member 3 (HACA small nucleolar RNPs) (NOLA3), mRNA. /FEA=mRNA /GEN=NOLA3 /PROD=nucleolar protein family A, member 3 (HACA small nucleolar RNPs) /DB_XREF=gi:8923941 /UG=Hs.14317 nucleolar protein family A, member 3 (HACA small nucleolar RNPs) /FL=gb:AB043104.1 gb:NM_018648.1	NM_018648		NP_061118
217967_s_at	0.033381	gb:AF288391.1 /DEF=Homo sapiens C1orf24 mRNA, complete cds. /FEA=mRNA /PROD=C1orf24 /DB_XREF=gi:12620191 /UG=Hs.48778 niban protein /FL=gb:AB050477.1 gb:NM_022083.1 gb:AF288391.1	AF288391		NP_443198

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217969_at	0.041055	gb:NM_013265.2 /DEF=Homo sapiens chromosome 11 open reading frame2 (C11ORF2), mRNA. /FEA=mRNA /GEN=C11ORF2 /PROD=chromosome 11 open reading frame2 /DB_XREF=gi:8393008 /UG=Hs.5258 chromosome 11 open reading frame2 /FL=gb:AF024631.2 gb:NM_013265.2	NM_013265		NP_008917
217970_s_at	0.002313	gb:NM_015455.1 /DEF=Homo sapiens KIAA1194 protein (KIAA1194), mRNA. /FEA=mRNA /GEN=KIAA1194 /PROD=DKFZP564G1964 protein /DB_XREF=gi:7661613 /UG=Hs.5801 KIAA1194 protein /FL=gb:NM_015455.1	NM_015455		NP_056270
217971_at	1.26E-04	gb:NM_021970.1 /DEF=Homo sapiens MEK partner 1 (MP1), mRNA. /FEA=mRNA /GEN=MP1 /PROD=MEK partner 1 /DB_XREF=gi:11496276 /UG=Hs.6361 MEK partner 1 /FL=gb:AF130115.1 gb:NM_021970.1 gb:AF201947.1	NM_021970		NP_068805
217972_at	0.02145	gb:NM_017812.1 /DEF=Homo sapiens hypothetical protein FLJ20420 (FLJ20420), mRNA. /FEA=mRNA /GEN=FLJ20420 /PROD=hypothetical protein FLJ20420 /DB_XREF=gi:8923389 /UG=Hs.6693 hypothetical protein FLJ20420 /FL=gb:NM_017812.1	NM_017812		NP_060282
217977_at	0.00161	gb:NM_016332.1 /DEF=Homo sapiens selenoprotein X, 1 (SEPX1), mRNA. /FEA=mRNA /GEN=SEPX1 /PROD=selenoprotein X, 1 /DB_XREF=gi:7706510 /UG=Hs.279623 selenoprotein X, 1 /FL=gb:AF187272.1 gb:BC003127.1 gb:AF166124.1 gb:NM_016332.1	NM_016332		NP_057416

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217978_s_at	7.58E-04	gb:NM_017582.1 /DEF=Homo sapiens NICE-5 protein (HSA243666), mRNA. /FEA=mRNA /GEN=HSA243666 /PROD=NICE-5 protein /DB_XREF=gi:8923743 /UG=Hs.284233 NICE-5 protein /FL=gb:AF116721.1 gb:NM_017582.1	NM_017582		NP_060052
217980_s_at	0.005852	gb:NM_017840.1 /DEF=Homo sapiens hypothetical protein FLJ20484 (FLJ20484), mRNA. /FEA=mRNA /GEN=FLJ20484 /PROD=hypothetical protein FLJ20484 /DB_XREF=gi:8923447 /UG=Hs.5080 hypothetical protein FLJ20484 /FL=gb:AF275806.1 gb:BC001040.1 gb:NM_017840.1 gb:AF183428.1	NM_017840		NP_060310
217981_s_at	0.004341	gb:NM_012192.1 /DEF=Homo sapiens fracture callus 1 (rat) homolog (FXC1), mRNA. /FEA=mRNA /GEN=FXC1 /PROD=fracture callus 1 (rat) homolog /DB_XREF=gi:6912381 /UG=Hs.54943 fracture callus 1 (rat) homolog /FL=gb:AF150105.1 gb:AF152355.1 gb:NM_012192.1 gb:AF183415.1	NM_012192		NP_036324
217983_s_at	0.007779	gb:NM_003730.2 /DEF=Homo sapiens ribonuclease 6 precursor (RNASE6PL), mRNA. /FEA=mRNA /GEN=RNASE6PL /PROD=ribonuclease 6 precursor /DB_XREF=gi:5231227 /UG=Hs.8297 ribonuclease 6 precursor /FL=gb:BC001660.1 gb:BC001819.1 gb:U85625.2 gb:NM_003730.2	NM_003730		NP_003721

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217986_s_at	2.05E-04	gb:NM_013448.1 /DEF=Homo sapiens bromodomain adjacent to zinc finger domain, 1A (BAZ1A), mRNA. /FEA=mRNA /GEN=BAZ1A /PROD=bromodomain adjacent to zinc finger domain, 1A /DB_XREF=gi:7304918 /UG=Hs.8858 bromodomain adjacent to zinc finger domain, 1A /FL=gb:AB032252.1 gb:NM_013448.1	NM_013448		NP_038476
217989_at	0.010205	gb:NM_016245.1 /DEF=Homo sapiens retinal short-chain dehydrogenasereductase retSDR2 (LOC51170), mRNA. /FEA=mRNA /GEN=LOC51170 /PROD=retinal short-chain dehydrogenasereductaseretSDR2 /DB_XREF=gi:7705904 /UG=Hs.12150 retinal short-chain dehydrogenasereductase retSDR2 /FL=gb:AF126780.1 gb:NM_016245.1	NM_016245		NP_057329
217990_at	7.58E-04	gb:NM_016576.1 /DEF=Homo sapiens GMPR2 for guanosine monophosphate reductase isolog (LOC51292), mRNA. /FEA=mRNA /GEN=LOC51292 /PROD=GMPR2 for guanosine monophosphate reductaseisolog /DB_XREF=gi:7706108 /UG=Hs.234546 GMPR2 for guanosine monophosphate reductase isolog /FL=gb:AB032903.1 gb:NM_016576.1 gb:AF135159.1	NM_016576		NP_057660
217995_at	1.26E-04	gb:NM_021199.1 /DEF=Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA. /FEA=mRNA /GEN=CGI-44 /PROD=CGI-44 protein; sulfide dehydrogenase like(yeast) /DB_XREF=gi:10864010 /UG=Hs.8185 CGI-44 protein; sulfide dehydrogenase like (yeast) /FL=gb:NM_021199.1 gb:AF151802.1 gb:AF118085.1	NM_021199		NP_067022

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218004_at	0.016934	gb:NM_018045.1 /DEF=Homo sapiens hypothetical protein FLJ10276 (FLJ10276), mRNA. /FEA=mRNA /GEN=FLJ10276 /PROD=hypothetical protein FLJ10276 /DB_XREF=gi:8922323 /UG=Hs.6937 hypothetical protein FLJ10276 /FL=gb:NM_018045.1	NM_018045		NP_060515
218006_s_at	0.001618	gb:NM_006963.1 /DEF=Homo sapiens zinc finger protein 22 (KOX 15) (ZNF22), mRNA. /FEA=mRNA /GEN=ZNF22 /PROD=zinc finger protein 22 (KOX 15) /DB_XREF=gi:5902159 /UG=Hs.108642 zinc finger protein 22 (KOX 15) /FL=gb:NM_006963.1	NM_006963		NP_008894
218008_at	7.58E-04	gb:NM_017994.1 /DEF=Homo sapiens hypothetical protein FLJ10099 (FLJ10099), mRNA. /FEA=mRNA /GEN=FLJ10099 /PROD=hypothetical protein FLJ10099 /DB_XREF=gi:8922228 /UG=Hs.278619 hypothetical protein FLJ10099 /FL=gb:NM_017994.1	NM_017994		NP_060464
218011_at	1.26E-04	gb:NM_024292.1 /DEF=Homo sapiens ubiquitin-like 5 (UBL5), mRNA. /FEA=mRNA /GEN=UBL5 /PROD=ubiquitin-like 5 /DB_XREF=gi:13236509 /UG=Hs.109701 ubiquitin-like 5 /FL=gb:AF313915.1 gb:NM_024292.1	NM_024292		NP_077268
218013_x_at	0.034221	gb:NM_016221.1 /DEF=Homo sapiens dynactin p62 subunit (LOC51164), mRNA. /FEA=mRNA /GEN=LOC51164 /PROD=dynactin p62 subunit /DB_XREF=gi:7705892 /UG=Hs.180952 dynactin 4 (p62) /FL=gb:AF195120.1 gb:NM_016221.1	NM_016221		NP_057305

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218017_s_at	0.026891	gb:NM_025070.1 /DEF=Homo sapiens hypothetical protein FLJ22242 (FLJ22242), mRNA. /FEA=mRNA /GEN=FLJ22242 /PROD=hypothetical protein FLJ22242 /DB_XREF=gi:13376612 /UG=Hs.288057 hypothetical protein FLJ22242 /FL=gb:NM_025070.1	NM_025070		NP_079346
218018_at	0.021448	Consensus includes gb:AW449022 /FEA=EST /DB_XREF=gi:6989798 /DB_XREF=est:UI-H-BI3-ake-f-11-0 UI.s1 /CLONE=IMAGE:2734220 /UG=Hs.4746 hypothetical protein FLJ21324 /FL=gb:NM_021941.1 gb:BC003651.1	NM_021941		NP_068760
218019_s_at	0.013223	gb:NM_021941.1 /DEF=Homo sapiens hypothetical protein FLJ21324 (FLJ21324), mRNA. /FEA=mRNA /GEN=FLJ21324 /PROD=hypothetical protein FLJ21324 /DB_XREF=gi:11345479 /UG=Hs.4746 hypothetical protein FLJ21324 /FL=gb:NM_021941.1 gb:BC003651.1	NM_021941		NP_068760
218023_s_at	5.03E-04	gb:NM_016605.1 /DEF=Homo sapiens putative nuclear protein (LOC51307), mRNA. /FEA=mRNA /GEN=LOC51307 /PROD=putative nuclear protein /DB_XREF=gi:7706138 /UG=Hs.102469 putative nuclear protein /FL=gb:AF251040.1 gb:NM_016605.1	NM_016605		NP_057689

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218030_at	1.26E-04	gb:NM_014030.1 /DEF=Homo sapiens G protein-coupled receptor kinase-interactor 1 (GIT1), mRNA. /FEA=mRNA /GEN=GIT1 /PROD=G protein-coupled receptor kinase-interactor 1 /DB_XREF=gi:7661711 /UG=Hs.318339 G protein-coupled receptor kinase-interactor 1 /FL=gb:AF124490.1 gb:NM_014030.1	NM_014030		NP_054749
218032_at	0.007779	gb:AF070673.1 /DEF=Homo sapiens stannin mRNA, complete cds. /FEA=mRNA /PROD=stannin /DB_XREF=gi:3978241 /UG=Hs.76691 stannin /FL=gb:AF030196.1 gb:AF070673.1 gb:NM_003498.1 gb:AL161976.1	AF070673		NP_003489
218034_at	0.005852	gb:NM_016068.1 /DEF=Homo sapiens CGI-135 protein (LOC51024), mRNA. /FEA=mRNA /GEN=LOC51024 /PROD=CGI-135 protein /DB_XREF=gi:7705631 /UG=Hs.84344 CGI-135 protein /FL=gb:BC003540.1 gb:AF151893.1 gb:NM_016068.1	NM_016068		NP_057152
218037_at	0.00161	gb:NM_024293.1 /DEF=Homo sapiens hypothetical protein MGC3035 (MGC3035), mRNA. /FEA=mRNA /GEN=MGC3035 /PROD=hypothetical protein MGC3035 /DB_XREF=gi:13236511 /UG=Hs.22412 hypothetical protein MGC3035 /FL=gb:AL136758.1 gb:BC002420.1 gb:NM_024293.1	NM_024293		NP_077269

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218039_at	0.00161	gb:NM_016359.1 /DEF=Homo sapiens clone HQ0310 PRO0310p1 (LOC51203), mRNA. /FEA=mRNA /GEN=LOC51203 /PROD=clone HQ0310 PRO0310p1 /DB_XREF=gi:7705950 /UG=Hs.279905 clone HQ0310 PRO0310p1 /FL=gb:AF305711.1 gb:BC001308.1 gb:AF290612.1 gb:AF090915.1 gb:NM_016359.1	NM_016359		NP_060924
218041_x_at	1.26E-04	gb:NM_018573.1 /DEF=Homo sapiens hypothetical protein PRO1068 (PRO1068), mRNA. /FEA=mRNA /GEN=PRO1068 /PROD=hypothetical protein PRO1068 /DB_XREF=gi:8924006 /UG=Hs.321158 hypothetical protein PRO1068 /FL=gb:AF116620.1 gb:NM_018573.1	NM_018573		
218043_s_at	0.00228	gb:NM_022461.1 /DEF=Homo sapiens hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 (FLJ21939), mRNA. /FEA=mRNA /GEN=FLJ21939 /PROD=hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 /DB_XREF=gi:11968002 /UG=Hs.164478 hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 /FL=gb:NM_022461.1	NM_022461		NP_071906
218047_at	5.03E-04	gb:NM_024586.1 /DEF=Homo sapiens hypothetical protein FLJ12492 (FLJ12492), mRNA. /FEA=mRNA /GEN=FLJ12492 /PROD=hypothetical protein FLJ12492 /DB_XREF=gi:13375771 /UG=Hs.21938 hypothetical protein FLJ12492 /FL=gb:NM_024586.1	NM_024586		NP_683707

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218049_s_at	0.02141	gb:NM_014078.1 /DEF=Homo sapiens L13 protein (L13), mRNA. /FEA=mRNA /GEN=L13 /PROD=L13 protein /DB_XREF=gi:7662495 /UG=Hs.43946 L13 protein /FL=gb:AF112214.1 gb:NM_014078.1	NM_014078		NP_054797
218057_x_at	0.003171	gb:NM_006067.1 /DEF=Homo sapiens neighbor of COX4 (NOC4), mRNA. /FEA=mRNA /GEN=NOC4 /PROD=neighbor of COX4 /DB_XREF=gi:5174614 /UG=Hs.173162 neighbor of COX4 /FL=gb:BC001472.1 gb:AF005888.1 gb:NM_006067.1	NM_006067		NP_006058
218066_at	0.016934	gb:NM_006598.1 /DEF=Homo sapiens solute carrier family 12 (potassiumchloride transporters), member 7 (SLC12A7), mRNA. /FEA=mRNA /GEN=SLC12A7 /PROD=solute carrier family 12 (potassiumchloridetransporters), member 7 /DB_XREF=gi:5730042 /UG=Hs.172613 solute carrier family 12 (potassiumchloride transporters), member 7 /FL=gb:AF105365.1 gb:NM_006598.1	NM_006598		NP_006589
218068_s_at	0.026891	gb:NM_024836.1 /DEF=Homo sapiens hypothetical protein FLJ22301 (FLJ22301), mRNA. /FEA=mRNA /GEN=FLJ22301 /PROD=hypothetical protein FLJ22301 /DB_XREF=gi:13376246 /UG=Hs.181406 hypothetical protein FLJ22301 /FL=gb:NM_024836.1	NM_024836		NP_079112
218071_s_at	7.58E-04	gb:NM_014160.2 /DEF=Homo sapiens HSPC070 protein (HSPC070), mRNA. /FEA=mRNA /GEN=HSPC070 /PROD=HSPC070 protein /DB_XREF=gi:8850222 /UG=Hs.279474 HSPC070 protein /FL=gb:AF302084.1 gb:AF161555.2 gb:NM_014160.2	NM_014160		NP_054879

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218077_s_at	0.00476	601063935F1 NIH_MGC_10 Homo sapiens cDNA clone IMAGE:3450249 5', mRNA sequence.	BE542551		
218078_s_at	1.26E-04	gb:NM_016598.1 /DEF=Homo sapiens DHHC1 protein (LOC51304), mRNA. /FEA=mRNA /GEN=LOC51304 /PROD=DHHC1 protein /DB_XREF=gi:7706132 /UG=Hs.14896 DHHC1 protein /FL=gb:AF247703.1 gb:NM_016598.1	NM_016598		NP_057682
218080_x_at	0.034317	gb:NM_007051.1 /DEF=Homo sapiens Fas (TNFRSF6) associated factor 1 (FAF1), mRNA. /FEA=mRNA /GEN=FAF1 /PROD=Fas (TNFRSF6) associated factor 1 /DB_XREF=gi:5901947 /UG=Hs.25821 Fas (TNFRSF6) associated factor 1 /FL=gb:BC004970.1 gb:AF132938.1 gb:AF106798.1 gb:NM_007051.1	NM_007051		NP_572051
218085_at	0.00161	gb:NM_015961.1 /DEF=Homo sapiens CGI-34 protein (LOC51612), mRNA. /FEA=mRNA /GEN=LOC51612 /PROD=CGI-34 protein /DB_XREF=gi:7706273 /UG=Hs.36237 CGI-34 protein /FL=gb:AF275810.1 gb:AF132968.1 gb:NM_015961.1	NM_015961		
218091_at	1.26E-04	HIV-1 Rev binding protein	AI989512	Hs.171545	NP_004495
218092_s_at	1.26E-04	gb:NM_004504.2 /DEF=Homo sapiens HIV-1 Rev binding protein (HRB), mRNA. /FEA=mRNA /GEN=HRB /PROD=HIV-1 Rev binding protein /DB_XREF=gi:7262381 /UG=Hs.171545 HIV-1 Rev binding protein /FL=gb:NM_004504.2	NM_004504		NP_004495

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218093_s_at	7.58E-04	gb:NM_017664.1 /DEF=Homo sapiens hypothetical protein FLJ20093 (FLJ20093), mRNA. /FEA=mRNA /GEN=FLJ20093 /PROD=hypothetical protein FLJ20093 /DB_XREF=gi:8923103 /UG=Hs.172572 hypothetical protein FLJ20093 /FL=gb:NM_017664.1	NM_017664		NP_060134
218095_s_at	0.010205	gb:NM_018475.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HTMP (LOC55858), mRNA. /FEA=mRNA /GEN=LOC55858 /PROD=uncharacterized hypothalamus protein HTMP /DB_XREF=gi:8923860 /UG=Hs.236510 uncharacterized hypothalamus protein HTMP /FL=gb:BC003545.1 gb:AF220188.1 gb:NM_018475.1 gb:AF183409.1	NM_018475		NP_060945
218097_s_at	0.003171	gb:NM_024040.1 /DEF=Homo sapiens hypothetical protein MGC2491 (MGC2491), mRNA. /FEA=mRNA /GEN=MGC2491 /PROD=hypothetical protein MGC2491 /DB_XREF=gi:13128995 /UG=Hs.11270 hypothetical protein MGC2491 /FL=gb:BC000262.1 gb:NM_024040.1	NM_024040		NP_076945
218099_at	0.041973	gb:NM_018469.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HT008 (HT008), mRNA. /FEA=mRNA /GEN=HT008 /PROD=uncharacterized hypothalamus protein HT008 /DB_XREF=gi:8923803 /UG=Hs.16206 uncharacterized hypothalamus protein HT008 /FL=gb:AF220182.1 gb:NM_018469.1	NM_018469		NP_060939

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218101_s_at	3.26E-04	gb:NM_004549.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) (NDUFC2), mRNA. /FEA=mRNA /GEN=NDUFC2 /PROD=NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) /DB_XREF=gi:4758783 /UG=Hs.193313 NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) /FL=gb:AF087659.1 gb:AF070652.1 gb:NM_004549.1	NM_004549		NP_004540
218102_at	0.00161	gb:NM_015954.1 /DEF=Homo sapiens CGI-26 protein (LOC51071), mRNA. /FEA=mRNA /GEN=LOC51071 /PROD=CGI-26 protein /DB_XREF=gi:7705717 /UG=Hs.24332 CGI-26 protein /FL=gb:AF132960.1 gb:NM_015954.1	NM_015954		NP_057038
218107_at	2.05E-04	gb:NM_025160.1 /DEF=Homo sapiens hypothetical protein FLJ21016 (FLJ21016), mRNA. /FEA=mRNA /GEN=FLJ21016 /PROD=hypothetical protein FLJ21016 /DB_XREF=gi:13376760 /UG=Hs.289069 hypothetical protein FLJ21016 /FL=gb:NM_025160.1	NM_025160		NP_079436
218108_at	2.05E-04	gb:NM_018108.1 /DEF=Homo sapiens hypothetical protein FLJ10483 (FLJ10483), mRNA. /FEA=mRNA /GEN=FLJ10483 /PROD=hypothetical protein FLJ10483 /DB_XREF=gi:8922451 /UG=Hs.6877 hypothetical protein FLJ10483 /FL=gb:NM_018108.1	NM_018108		NP_786924

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218109_s_at	1.26E-04	gb:NM_022736.1 /DEF=Homo sapiens hypothetical protein FLJ14153 (FLJ14153), mRNA. /FEA=mRNA /GEN=FLJ14153 /PROD=hypothetical protein FLJ14153 /DB_XREF=gi:12232392 /UG=Hs.7503 hypothetical protein FLJ14153 /FL=gb:NM_022736.1 gb:AB014732.1	NM_022736		NP_073573
218111_s_at	0.002313	gb:NM_018686.1 /DEF=Homo sapiens CMP-N-acetylneuraminic acid synthase (LOC55907), mRNA. /FEA=mRNA /GEN=LOC55907 /PROD=CMP-N-acetylneuraminic acid synthase /DB_XREF=gi:8923899 /UG=Hs.12492 CMP-N-acetylneuraminic acid synthase /FL=gb:AF271388.1 gb:NM_018686.1	NM_018686		NP_061156
218113_at	5.03E-04	gb:NM_013390.1 /DEF=Homo sapiens transmembrane protein 2 (TMEM2), mRNA. /FEA=mRNA /GEN=TMEM2 /PROD=transmembrane protein 2 /DB_XREF=gi:7019554 /UG=Hs.160417 transmembrane protein 2 /FL=gb:AF137030.1 gb:NM_013390.1	NM_013390		NP_037522
218116_at	0.026891	gb:NM_016520.1 /DEF=Homo sapiens hepatocellular carcinoma-associated antigen 59 (LOC51759), mRNA. /FEA=mRNA /GEN=LOC51759 /PROD=hepatocellular carcinoma-associated antigen 59 /DB_XREF=gi:7706556 /UG=Hs.278429 hepatocellular carcinoma-associated antigen 59 /FL=gb:AF151054.1 gb:AF218421.1 gb:NM_016482.1 gb:NM_016520.1	NM_016520		NP_057604

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218117_at	0.00228	gb:NM_014248.1 /DEF=Homo sapiens ring-box 1 (RBX1), mRNA. /FEA=mRNA /GEN=RBX1 /PROD=ring-box 1 /DB_XREF=gi:7657507 /UG=Hs.279919 ring-box 1 /FL=gb:BC001466.1 gb:AF140598.1 gb:AF142059.1 gb:NM_014248.1	NM_014248		NP_055063
218118_s_at	0.010061	gb:NM_006327.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 23 (yeast) homolog (TIM23), mRNA. /FEA=mRNA /GEN=TIM23 /PROD=translocase of inner mitochondrial membrane 23(yeast) homolog /DB_XREF=gi:5454121 /UG=Hs.283684 translocase of inner mitochondrial membrane 23 (yeast) homolog /FL=gb:AF030162.1 gb:NM_006327.1	NM_006327		NP_006318
218119_at	0.033381	gb:NM_006327.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 23 (yeast) homolog (TIM23), mRNA. /FEA=mRNA /GEN=TIM23 /PROD=translocase of inner mitochondrial membrane 23(yeast) homolog /DB_XREF=gi:5454121 /UG=Hs.283684 translocase of inner mitochondrial membrane 23 (yeast) homolog /FL=gb:AF030162.1 gb:NM_006327.1	NM_006327		NP_006318
218122_s_at	0.013088	gb:NM_021627.1 /DEF=Homo sapiens sentrin-specific protease (SEN2), mRNA. /FEA=mRNA /GEN=SEN2 /PROD=sentrin-specific protease /DB_XREF=gi:11055993 /UG=Hs.3355 sentrin-specific protease /FL=gb:AF151697.2 gb:NM_021627.1	NM_021627		NP_067640

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218124_at	7.58E-04	gb:NM_017750.1 /DEF=Homo sapiens hypothetical protein FLJ20296 (FLJ20296), mRNA. /FEA=mRNA /GEN=FLJ20296 /PROD=hypothetical protein FLJ20296 /DB_XREF=gi:8923274 /UG=Hs.6603 hypothetical protein FLJ20296 /FL=gb:NM_017750.1	NM_017750		NP_060220
218129_s_at	0.001078	gb:NM_006166.2 /DEF=Homo sapiens nuclear transcription factor Y, beta (NFYB), mRNA. /FEA=mRNA /GEN=NFYB /PROD=nuclear transcription factor Y, beta /DB_XREF=gi:11496976 /UG=Hs.84928 nuclear transcription factor Y, beta /FL=gb:NM_006166.2 gb:BC005316.1 gb:BC005317.1 gb:L06145.1	NM_006166		NP_006157
218130_at	1.26E-04	gb:NM_024510.1 /DEF=Homo sapiens hypothetical protein MGC4368 (MGC4368), mRNA. /FEA=mRNA /GEN=MGC4368 /PROD=hypothetical protein MGC4368 /DB_XREF=gi:13375647 /UG=Hs.9732 hypothetical protein MGC4368 /FL=gb:BC003595.1 gb:BC004171.1 gb:NM_024510.1	NM_024510		NP_078786
218131_s_at	0.004341	gb:NM_017660.1 /DEF=Homo sapiens hypothetical protein FLJ20085 (FLJ20085), mRNA. /FEA=mRNA /GEN=FLJ20085 /PROD=hypothetical protein FLJ20085 /DB_XREF=gi:8923093 /UG=Hs.118964 hypothetical protein FLJ20085 /FL=gb:NM_017660.1	NM_017660		NP_060130

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218134_s_at	0.010205	gb:NM_018047.1 /DEF=Homo sapiens hypothetical protein FLJ10290 (FLJ10290), mRNA. /FEA=mRNA /GEN=FLJ10290 /PROD=hypothetical protein FLJ10290 /DB_XREF=gi:8922327 /UG=Hs.25516 hypothetical protein FLJ10290 /FL=gb:AL136933.1 gb:BC003402.1 gb:NM_018047.1	NM_018047		NP_060517
218135_at	0.00228	gb:NM_016570.1 /DEF=Homo sapiens CDA14 (LOC51290), mRNA. /FEA=mRNA /GEN=LOC51290 /PROD=CDA14 /DB_XREF=gi:7706104 /UG=Hs.26813 CDA14 /FL=gb:BC000887.1 gb:AF216751.1 gb:NM_016570.1 gb:AF183410.1	NM_016570		NP_057654
218137_s_at	5.03E-04	gb:NM_021940.1 /DEF=Homo sapiens hypothetical protein FLJ13159 (FLJ13159), mRNA. /FEA=mRNA /GEN=FLJ13159 /PROD=hypothetical protein FLJ13159 /DB_XREF=gi:11345477 /UG=Hs.4273 hypothetical protein FLJ13159 /FL=gb:NM_021940.1	NM_021940		NP_068759
218138_at	0.005852	gb:NM_018848.1 /DEF=Homo sapiens McKusick-Kaufman syndrome (MKKS), mRNA. /FEA=mRNA /GEN=MKKS /PROD=McKusick-Kaufman syndrome protein /DB_XREF=gi:9055271 /UG=Hs.46743 McKusick-Kaufman syndrome /FL=gb:AF275813.1 gb:AF221992.1 gb:AF221993.1 gb:NM_018848.1	NM_018848		NP_740754

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218139_s_at	0.010205	gb:NM_018229.1 /DEF=Homo sapiens hypothetical protein FLJ10813 (FLJ10813), mRNA. /FEA=mRNA /GEN=FLJ10813 /PROD=hypothetical protein FLJ10813 /DB_XREF=gi:8922687 /UG=Hs.106210 hypothetical protein FLJ10813 /FL=gb:AL136685.1 gb:NM_018229.1	NM_018229		NP_060699
218142_s_at	0.00161	gb:NM_016302.1 /DEF=Homo sapiens protein x 0001 (LOC51185), mRNA. /FEA=mRNA /GEN=LOC51185 /PROD=protein x 0001 /DB_XREF=gi:10047097 /UG=Hs.18925 protein x 0001 /FL=gb:NM_016302.1 gb:AF117230.1	NM_016302		NP_057386
218157_x_at	0.00161	gb:NM_020239.2 /DEF=Homo sapiens small protein effector 1 of Cdc42 (SPEC1), mRNA. /FEA=mRNA /GEN=SPEC1 /PROD=small protein effector 1 of Cdc42 /DB_XREF=gi:12965169 /UG=Hs.22065 small protein effector 1 of Cdc42 /FL=gb:AF187845.2 gb:NM_020239.2	NM_020239		NP_064624
218158_s_at	0.036254	gb:NM_012096.1 /DEF=Homo sapiens adaptor protein containing pH domain, PTB domain and leucine zipper motif (APPL), mRNA. /FEA=mRNA /GEN=APPL /PROD=adaptor protein containing pH domain, PTB domain and leucine zipper motif /DB_XREF=gi:6912241 /UG=Hs.27413 adaptor protein containing pH domain, PTB domain and leucine zipper motif /FL=gb:AF169797.1 gb:NM_012096.1	NM_012096		NP_036228

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218159_at	0.007258	gb:NM_023935.1 /DEF=Homo sapiens hypothetical protein MGC2592 (MGC2592), mRNA. /FEA=mRNA /GEN=MGC2592 /PROD=hypothetical protein MGC2592 /DB_XREF=gi:13027601 /UG=Hs.28393 hypothetical protein MGC2592 /FL=gb:BC000643.1 gb:NM_023935.1	NM_023935		NP_076424
218163_at	2.05E-04	gb:NM_014060.1 /DEF=Homo sapiens MCT-1 protein (MCT-1), mRNA. /FEA=mRNA /GEN=MCT-1 /PROD=MCT-1 protein /DB_XREF=gi:7662501 /UG=Hs.102696 MCT-1 protein /FL=gb:BC001013.1 gb:AB034206.1 gb:NM_014060.1	NM_014060		NP_054779
218167_at	0.033381	gb:NM_016627.1 /DEF=Homo sapiens hypothetical protein (LOC51321), mRNA. /FEA=mRNA /GEN=LOC51321 /PROD=hypothetical protein /DB_XREF=gi:7706167 /UG=Hs.268122 hypothetical protein /FL=gb:AF208856.1 gb:NM_016627.1	NM_016627		NP_057711
218168_s_at	0.007779	gb:NM_020247.1 /DEF=Homo sapiens hypothetical protein, clone Telethon(Italy_B41)_Strait02270_F L142 (LOC56997), mRNA. /FEA=mRNA /GEN=LOC56997 /PROD=hypothetical protein, clone Telethon(Italy_B41)_Strait02270_FL142 /DB_XREF=gi:9910387 /UG=Hs.273186 hypothetical protein, clone Telethon(Italy_B41)_Strait02270_F L142 /FL=gb:BC005171.1 gb:NM_020247.1	NM_020247		NP_064632

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218171_at	2.05E-04	gb:AF195514.1 /DEF=Homo sapiens VPS4-2 ATPase (VPS42) mRNA, complete cds. /FEA=mRNA /GEN=VPS42 /PROD=VPS4-2 ATPase /DB_XREF=gi:11225484 /UG=Hs.126550 suppressor of K+ transport defect 1 /FL=gb:AF195514.1 gb:AF038960.1 gb:NM_004869.1	AF195514		NP_004860
218177_at	0.016934	CHMP1.5 protein	AA293502	Hs.380932	NP_065145
218178_s_at	0.02145	gb:NM_020412.1 /DEF=Homo sapiens CHMP1.5 protein (CHMP1.5), mRNA. /FEA=mRNA /GEN=CHMP1.5 /PROD=CHMP1.5 protein /DB_XREF=gi:9966900 /UG=Hs.42733 CHMP1.5 protein /FL=gb:AF281064.1 gb:NM_020412.1	NM_020412		NP_065145
218179_s_at	0.02145	gb:NM_021942.1 /DEF=Homo sapiens hypothetical protein FLJ12716 (FLJ12716), mRNA. /FEA=mRNA /GEN=FLJ12716 /PROD=hypothetical protein FLJ12716 /DB_XREF=gi:11345481 /UG=Hs.5354 hypothetical protein FLJ12716 /FL=gb:NM_021942.1 gb:AL136752.1	NM_021942		NP_068761
218181_s_at	1.26E-04	gb:NM_017792.1 /DEF=Homo sapiens hypothetical protein FLJ20373 (FLJ20373), mRNA. /FEA=mRNA /GEN=FLJ20373 /PROD=hypothetical protein FLJ20373 /DB_XREF=gi:8923351 /UG=Hs.6631 hypothetical protein FLJ20373 /FL=gb:NM_017792.1	NM_017792		NP_060262
218185_s_at	0.013223	gb:NM_018120.1 /DEF=Homo sapiens hypothetical protein FLJ10511 (FLJ10511), mRNA. /FEA=mRNA /GEN=FLJ10511 /PROD=hypothetical protein FLJ10511 /DB_XREF=gi:8922478 /UG=Hs.106768 hypothetical protein FLJ10511 /FL=gb:NM_018120.1	NM_018120		NP_060590

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218187_s_at	0.02145	gb:NM_023080.1 /DEF=Homo sapiens hypothetical protein FLJ20989 (FLJ20989), mRNA. /FEA=mRNA /GEN=FLJ20989 /PROD=hypothetical protein FLJ20989 /DB_XREF=gi:12751496 /UG=Hs.169615 hypothetical protein FLJ20989 /FL=gb:NM_023080.1	NM_023080		NP_075568
218189_s_at	3.26E-04	gb:NM_018946.2 /DEF=Homo sapiens N-acetylneuraminic acid phosphate synthase; sialic acid synthase (SAS), mRNA. /FEA=mRNA /GEN=SAS /PROD=N-acetylneuraminic acid phosphate synthase /DB_XREF=gi:12056472 /UG=Hs.274424 N-acetylneuraminic acid phosphate synthase; sialic acid synthase /FL=gb:NM_018946.2 gb:BC000008.1 gb:AF257466.1	NM_018946		NP_061819
218191_s_at	0.003171	gb:NM_018368.1 /DEF=Homo sapiens hypothetical protein FLJ11240 (FLJ11240), mRNA. /FEA=mRNA /GEN=FLJ11240 /PROD=hypothetical protein FLJ11240 /DB_XREF=gi:8922955 /UG=Hs.301308 hypothetical protein FLJ11240 /FL=gb:AF113224.1 gb:AF211480.1 gb:NM_018368.1	NM_018368		NP_060838
218195_at	1.26E-04	gb:NM_024573.1 /DEF=Homo sapiens hypothetical protein FLJ12910 (FLJ12910), mRNA. /FEA=mRNA /GEN=FLJ12910 /PROD=hypothetical protein FLJ12910 /DB_XREF=gi:13375745 /UG=Hs.15929 hypothetical protein FLJ12910 /FL=gb:NM_024573.1	NM_024573		NP_078849

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218196_at	3.26E-04	gb:NM_014028.1 /DEF=Homo sapiens HSPC019 protein (HSPC019), mRNA. /FEA=mRNA /GEN=HSPC019 /PROD=HSPC019 protein /DB_XREF=gi:7661737 /UG=Hs.163724 HSPC019 protein /FL=gb:AF077205.1 gb:NM_014028.1	NM_014028		NP_054747
218202_x_at	0.00161	gb:NM_022915.1 /DEF=Homo sapiens hypothetical protein FLJ12701 (FLJ12701), mRNA. /FEA=mRNA /GEN=FLJ12701 /PROD=hypothetical protein FLJ12701 /DB_XREF=gi:12597660 /UG=Hs.203559 hypothetical protein FLJ12701 /FL=gb:NM_022915.1	NM_022915		NP_075066
218208_at	3.26E-04	gb:NM_025078.1 /DEF=Homo sapiens hypothetical protein FLJ22378 (FLJ22378), mRNA. /FEA=mRNA /GEN=FLJ22378 /PROD=hypothetical protein FLJ22378 /DB_XREF=gi:13376629 /UG=Hs.288284 hypothetical protein FLJ22378 /FL=gb:NM_025078.1	NM_025078		NP_079354
218209_s_at	3.26E-04	gb:NM_018170.1 /DEF=Homo sapiens hypothetical protein FLJ10656 (FLJ10656), mRNA. /FEA=mRNA /GEN=FLJ10656 /PROD=hypothetical protein FLJ10656 /DB_XREF=gi:8922574 /UG=Hs.300906 hypothetical protein FLJ10656 /FL=gb:NM_018170.1	NM_018170		NP_060640

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218213_s_at	0.00161	gb:NM_014206.1 /DEF=Homo sapiens chromosome 11 open reading frame 10 (C11orf10), mRNA. /FEA=mRNA /GEN=C11orf10 /PROD=chromosome 11 open reading frame 10 /DB_XREF=gi:7656933 /UG=Hs.90918 chromosome 11 open reading frame 10 /FL=gb:BC002750.1 gb:AF070661.1 gb:AF086763.1 gb:NM_014206.1	NM_014206		NP_055021
218217_at	1.26E-04	gb:NM_021626.1 /DEF=Homo sapiens serine carboxypeptidase 1 precursor protein (HSCP1), mRNA. /FEA=mRNA /GEN=HSCP1 /PROD=serine carboxypeptidase 1 precursor protein /DB_XREF=gi:11055991 /UG=Hs.106747 serine carboxypeptidase 1 precursor protein /FL=gb:AF282618.1 gb:NM_021626.1 gb:AF113214.1 gb:AF265441.1	NM_021626		NP_067639
218218_at	1.26E-04	gb:NM_018171.1 /DEF=Homo sapiens hypothetical protein FLJ10659 (FLJ10659), mRNA. /FEA=mRNA /GEN=FLJ10659 /PROD=hypothetical protein FLJ10659 /DB_XREF=gi:8922576 /UG=Hs.107882 hypothetical protein FLJ10659 /FL=gb:NM_018171.1	NM_018171		NP_060641
218221_at	1.26E-04	ESTs	AL042842	Hs.166172	NP_848514
218223_s_at	0.001116	gb:NM_016274.1 /DEF=Homo sapiens CK2 interacting protein 1; HQ0024c protein (LOC51177), mRNA. /FEA=mRNA /GEN=LOC51177 /PROD=CK2 interacting protein 1; HQ0024c protein /DB_XREF=gi:7705918 /UG=Hs.173380 CK2 interacting protein 1; HQ0024c protein /FL=gb:AF291105.1 gb:AF073836.1 gb:NM_016274.1 gb:AF168676.1	NM_016274		NP_057358

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218226_s_at	0.010205	gb:NM_004547.2 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15) (NDUFB4), mRNA. /FEA=mRNA /GEN=NDUFB4 /PROD=NADH dehydrogenase (ubiquinone) 1 betasubcomplex, 4 (15kD, B15) /DB_XREF=gi:6041668 /UG=Hs.227750 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15) /FL=gb:BC000855.1 gb:AF044957.1 gb:NM_004547.2	NM_004547		NP_004538
218228_s_at	0.005852	gb:NM_025235.1 /DEF=Homo sapiens tankyrase 2 (TNKL), mRNA. /FEA=mRNA /GEN=TNKL /PROD=tankyrase 2 /DB_XREF=gi:13376841 /UG=Hs.280776 tankyrase 2 /FL=gb:AF264912.1 gb:AF329696.1 gb:NM_025235.1 gb:AF342982.1	NM_025235		NP_079511
218237_s_at	0.001116	gb:NM_030674.1 /DEF=Homo sapiens amino acid transporter system A1 (ATA1), mRNA. /FEA=mRNA /GEN=ATA1 /PROD=amino acid transporter system A1 /DB_XREF=gi:13492978 /UG=Hs.18272 amino acid transporter system A1 /FL=gb:AF271070.1 gb:NM_030674.1	NM_030674		NP_109599
218243_at	5.03E-04	gb:NM_025158.1 /DEF=Homo sapiens hypothetical protein FLJ22251 (FLJ22251), mRNA. /FEA=mRNA /GEN=FLJ22251 /PROD=hypothetical protein FLJ22251 /DB_XREF=gi:13376756 /UG=Hs.289064 hypothetical protein FLJ22251 /FL=gb:NM_025158.1	NM_025158		NP_079434

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218249_at	0.033381	gb:NM_022494.1 /DEF=Homo sapiens hypothetical protein FLJ21952 (FLJ21952), mRNA. /FEA=mRNA /GEN=FLJ21952 /PROD=hypothetical protein FLJ21952 /DB_XREF=gi:11968052 /UG=Hs.22353 hypothetical protein FLJ21952 /FL=gb:NM_022494.1	NM_022494		NP_071939
218250_s_at	0.004341	gb:NM_013354.2 /DEF=Homo sapiens CCR4-NOT transcription complex, subunit 7 (CNOT7), mRNA. /FEA=mRNA /GEN=CNOT7 /PROD=CCR4-NOT transcription complex, subunit 7 /DB_XREF=gi:10518495 /UG=Hs.226318 CCR4-NOT transcription complex, subunit 7 /FL=gb:NM_013354.2 gb:L46722.1	NM_013354		NP_473367
218251_at	2.05E-04	gb:NM_021242.1 /DEF=Homo sapiens hypothetical protein STRAIT11499 (STRAIT11499), mRNA. /FEA=mRNA /GEN=STRAIT11499 /PROD=hypothetical protein STRAIT11499 /DB_XREF=gi:10864050 /UG=Hs.236556 hypothetical protein STRAIT11499 /FL=gb:NM_021242.1	NM_021242		NP_067065
218252_at	3.26E-04	gb:NM_018204.1 /DEF=Homo sapiens cytoskeleton associated protein 2 (CKAP2), mRNA. /FEA=mRNA /GEN=CKAP2 /PROD=cytoskeleton associated protein 2 /DB_XREF=gi:8922641 /UG=Hs.24641 cytoskeleton associated protein 2 /FL=gb:AL136848.1 gb:NM_018204.1	NM_018204		NP_060674

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218253_s_at	0.028893	gb:NM_006893.1 /DEF=Homo sapiens ligatin (LGTN), mRNA. /FEA=mRNA /GEN=LGTN /PROD=ligatin /DB_XREF=gi:11386190 /UG=Hs.274151 ligatin /FL=gb:NM_006893.1 gb:AF220417.2	NM_006893		NP_008824
218254_s_at	2.05E-04	gb:NM_016103.1 /DEF=Homo sapiens GTP-binding protein Sara (LOC51128), mRNA. /FEA=mRNA /GEN=LOC51128 /PROD=GTP-binding protein Sara /DB_XREF=gi:7705826 /UG=Hs.279582 GTP-binding protein Sara /FL=gb:AF092130.1 gb:NM_016103.1	NM_016103		NP_057187
218255_s_at	0.041055	gb:NM_022452.1 /DEF=Homo sapiens hypothetical protein FLJ11618 (FLJ11618), mRNA. /FEA=mRNA /GEN=FLJ11618 /PROD=hypothetical protein FLJ11618 /DB_XREF=gi:11967986 /UG=Hs.77735 hypothetical protein FLJ11618 /FL=gb:NM_022452.1	NM_022452		NP_071897
218262_at	7.58E-04	gb:NM_022762.1 /DEF=Homo sapiens hypothetical protein FLJ22318 (FLJ22318), mRNA. /FEA=mRNA /GEN=FLJ22318 /PROD=hypothetical protein FLJ22318 /DB_XREF=gi:12232432 /UG=Hs.22753 hypothetical protein FLJ22318 /FL=gb:NM_022762.1	NM_022762		NP_073599
218268_at	0.013223	gb:NM_022771.1 /DEF=Homo sapiens hypothetical protein FLJ12085 (FLJ12085), mRNA. /FEA=mRNA /GEN=FLJ12085 /PROD=hypothetical protein FLJ12085 /DB_XREF=gi:12232450 /UG=Hs.48827 hypothetical protein FLJ12085 /FL=gb:NM_022771.1	NM_022771		NP_073608

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218271_s_at	3.26E-04	gb:NM_018622.1 /DEF=Homo sapiens presenilins associated rhomboid-like protein (PRO2207), mRNA. /FEA=mRNA /GEN=PRO2207 /PROD=presenilins associated rhomboid-like protein /DB_XREF=gi:8924133 /UG=Hs.13094 presenilins associated rhomboid-like protein /FL=gb:AF197937.1 gb:AF116692.1 gb:NM_018622.1	NM_018622		NP_061092
218273_s_at	7.58E-04	gb:NM_018444.1 /DEF=Homo sapiens pyruvate dehydrogenase phosphatase (PDP), mRNA. /FEA=mRNA /GEN=PDP /PROD=pyruvate dehydrogenase phosphatase /DB_XREF=gi:8923959 /UG=Hs.22265 pyruvate dehydrogenase phosphatase /FL=gb:AF155661.1 gb:NM_018444.1	NM_018444		NP_060914
218277_s_at	0.010205	gb:NM_024612.1 /DEF=Homo sapiens hypothetical protein FLJ22060 (FLJ22060), mRNA. /FEA=mRNA /GEN=FLJ22060 /PROD=hypothetical protein FLJ22060 /DB_XREF=gi:13375824 /UG=Hs.29403 hypothetical protein FLJ22060 /FL=gb:NM_024612.1	NM_024612		NP_078888
218280_x_at	5.03E-04	gb:NM_003516.1 /DEF=Homo sapiens H2A histone family, member O (H2AFO), mRNA. /FEA=mRNA /GEN=H2AFO /PROD=H2A histone family, member O /DB_XREF=gi:4504250 /UG=Hs.795 H2A histone family, member O /FL=gb:BC001629.1 gb:L19779.1 gb:NM_003516.1	NM_003516		NP_003507

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218286_s_at	0.003171	gb:NM_014245.1 /DEF=Homo sapiens ring finger protein 7 (RNF7), mRNA. /FEA=mRNA /GEN=RNF7 /PROD=ring finger protein 7 /DB_XREF=gi:7657521 /UG=Hs.14084 ring finger protein 7 /FL=gb:AF092878.1 gb:AF164679.1 gb:NM_014245.1	NM_014245		NP_055060
218291_at	7.58E-04	gb:NM_014017.1 /DEF=Homo sapiens HSPC003 protein (HSPC003), mRNA. /FEA=mRNA /GEN=HSPC003 /PROD=HSPC003 protein /DB_XREF=gi:7661727 /UG=Hs.25635 HSPC003 protein /FL=gb:AF070659.1 gb:NM_014017.1	NM_014017		NP_054736
218301_at	0.010205	gb:NM_018226.1 /DEF=Homo sapiens arginyl aminopeptidase (aminopeptidase B)-like 1 (RNPEPL1), mRNA. /FEA=mRNA /GEN=RNPEPL1 /PROD=arginyl aminopeptidase (aminopeptidase B)-like1 /DB_XREF=gi:8922680 /UG=Hs.5345 arginyl aminopeptidase (aminopeptidase B)-like 1 /FL=gb:AF300795.1 gb:NM_018226.1	NM_018226		NP_060696
218303_x_at	1.26E-04	gb:NM_016618.1 /DEF=Homo sapiens hypothetical protein (LOC51315), mRNA. /FEA=mRNA /GEN=LOC51315 /PROD=hypothetical protein /DB_XREF=gi:7706155 /UG=Hs.5721 hypothetical protein /FL=gb:AF208845.1 gb:AF217520.1 gb:NM_016618.1	NM_016618		NP_057702
218304_s_at	1.26E-04	gb:NM_022776.1 /DEF=Homo sapiens hypothetical protein FLJ13164 (FLJ13164), mRNA. /FEA=mRNA /GEN=FLJ13164 /PROD=hypothetical protein FLJ13164 /DB_XREF=gi:12232460 /UG=Hs.61260 hypothetical protein FLJ13164 /FL=gb:NM_022776.1	NM_022776		NP_073613

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218306_s_at	0.013223	gb:NM_003922.1 /DEF=Homo sapiens hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 (HERC1), mRNA. /FEA=mRNA /GEN=HERC1 /PROD=guanine nucleotide exchange factor p532 /DB_XREF=gi:4557025 /UG=Hs.76127 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 /FL=gb:U50078.1 gb:NM_003922.1	NM_003922		NP_003913
218308_at	0.02145	gb:NM_006342.1 /DEF=Homo sapiens transforming, acidic coiled-coil containing protein 3 (TACC3), mRNA. /FEA=mRNA /GEN=TACC3 /PROD=transforming acidic coiled-coil containing protein 3 /DB_XREF=gi:5454101 /UG=Hs.104019 transforming, acidic coiled-coil containing protein 3 /FL=gb:AF093543.1 gb:NM_006342.1	NM_006342		NP_006333
218310_at	2.05E-04	gb:NM_014504.1 /DEF=Homo sapiens putative Rab5 GDPGTP exchange factor homologue (RABEX5), mRNA. /FEA=mRNA /GEN=RABEX5 /PROD=putative Rab5 GDPGTP exchange factor homologue /DB_XREF=gi:7657495 /UG=Hs.187660 putative Rab5 GDPGTP exchange factor homologue /FL=gb:NM_014504.1	NM_014504		NP_055319

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218313_s_at	2.05E-04	gb:NM_017423.1 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7) (GALNT7), mRNA. /FEA=mRNA /GEN=GALNT7 /PROD=polypeptide N-acetylgalactosaminyltransferase 7 /DB_XREF=gi:8393408 /UG=Hs.246315 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7) /FL=gb:NM_017423.1	NM_017423		NP_059119
218318_s_at	0.04734	gb:NM_016231.1 /DEF=Homo sapiens nemo-like kinase (LOC51701), mRNA. /FEA=mRNA /GEN=LOC51701 /PROD=nemo-like kinase /DB_XREF=gi:7706444 /UG=Hs.3532 nemo-like kinase /FL=gb:AF180819.1 gb:AF197898.1 gb:NM_016231.1	NM_016231		NP_057315
218323_at	0.001116	gb:NM_018307.1 /DEF=Homo sapiens hypothetical protein FLJ11040 (FLJ11040), mRNA. /FEA=mRNA /GEN=FLJ11040 /PROD=hypothetical protein FLJ11040 /DB_XREF=gi:8922837 /UG=Hs.14202 hypothetical protein FLJ11040 /FL=gb:NM_018307.1	NM_018307		NP_060777
218332_at	8.94E-04	gb:NM_018476.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HBEX2 (HBEX2), mRNA. /FEA=mRNA /GEN=HBEX2 /PROD=uncharacterized hypothalamus protein HBEX2 /DB_XREF=gi:8923715 /UG=Hs.283719 uncharacterized hypothalamus protein HBEX2 /FL=gb:AF220189.1 gb:NM_018476.1 gb:AF183416.1 gb:AF237783.1	NM_018476		NP_060946

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218333_at	0.007779	gb:NM_016041.1 /DEF=Homo sapiens CGI-101 protein (LOC51009), mRNA. /FEA=mRNA /GEN=LOC51009 /PROD=CGI-101 protein /DB_XREF=gi:7705603 /UG=Hs.286131 CGI-101 protein /FL=gb:AF132289.1 gb:AF151859.1 gb:NM_016041.1 gb:AF242523.1	NM_016041		NP_057125
218334_at	0.00228	gb:NM_025075.1 /DEF=Homo sapiens hypothetical protein FLJ23445 (FLJ23445), mRNA. /FEA=mRNA /GEN=FLJ23445 /PROD=hypothetical protein FLJ23445 /DB_XREF=gi:13376622 /UG=Hs.288151 hypothetical protein FLJ23445 /FL=gb:NM_025075.1	NM_025075		NP_079351
218341_at	0.004341	gb:NM_024664.1 /DEF=Homo sapiens hypothetical protein FLJ11838 (FLJ11838), mRNA. /FEA=mRNA /GEN=FLJ11838 /PROD=hypothetical protein FLJ11838 /DB_XREF=gi:13375918 /UG=Hs.72531 hypothetical protein FLJ11838 /FL=gb:NM_024664.1	NM_024664		NP_078940
218343_s_at	0.026891	gb:NM_012086.1 /DEF=Homo sapiens general transcription factor IIIC, polypeptide 3 (102kD) (GTF3C3), mRNA. /FEA=mRNA /GEN=GTF3C3 /PROD=general transcription factor IIIC, polypeptide 3(102kD) /DB_XREF=gi:6912397 /UG=Hs.90847 general transcription factor IIIC, polypeptide 3 (102kD) /FL=gb:AF133123.1 gb:NM_012086.1	NM_012086		NP_036218

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218344_s_at	0.013223	gb:NM_018254.1 /DEF=Homo sapiens hypothetical protein FLJ10876 (FLJ10876), mRNA. /FEA=mRNA /GEN=FLJ10876 /PROD=hypothetical protein FLJ10876 /DB_XREF=gi:8922732 /UG=Hs.94042 hypothetical protein FLJ10876 /FL=gb:NM_018254.1	NM_018254		NP_060724
218345_at	0.02049	gb:NM_018487.1 /DEF=Homo sapiens hepatocellular carcinoma-associated antigen 112 (HCA112), mRNA. /FEA=mRNA /GEN=HCA112 /PROD=hepatocellular carcinoma-associated antigen 112 /DB_XREF=gi:8923717 /UG=Hs.12126 hepatocellular carcinoma-associated antigen 112 /FL=gb:AF258340.1 gb:NM_018487.1	NM_018487		NP_060957
218348_s_at	2.05E-04	gb:NM_014153.1 /DEF=Homo sapiens HSPC055 protein (HSPC055), mRNA. /FEA=mRNA /GEN=HSPC055 /PROD=HSPC055 protein /DB_XREF=gi:7661761 /UG=Hs.179898 HSPC055 protein /FL=gb:AF161540.1 gb:NM_014153.1	NM_014153		NP_054872
218350_s_at	0.002743	gb:NM_015895.1 /DEF=Homo sapiens geminin (LOC51053), mRNA. /FEA=mRNA /GEN=LOC51053 /PROD=geminin /DB_XREF=gi:7705681 /UG=Hs.234896 geminin /FL=gb:AF067855.1 gb:NM_015895.1	NM_015895		NP_056979
218351_at	0.026891	gb:NM_017845.1 /DEF=Homo sapiens hypothetical protein FLJ20502 (FLJ20502), mRNA. /FEA=mRNA /GEN=FLJ20502 /PROD=hypothetical protein FLJ20502 /DB_XREF=gi:8923457 /UG=Hs.23956 hypothetical protein FLJ20502 /FL=gb:AF182421.1 gb:NM_017845.1	NM_017845		NP_060315

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218352_at	3.26E-04	gb:NM_018191.1 /DEF=Homo sapiens hypothetical protein FLJ10716 (FLJ10716), mRNA. /FEA=mRNA /GEN=FLJ10716 /PROD=hypothetical protein FLJ10716 /DB_XREF=gi:8922616 /UG=Hs.24129 hypothetical protein FLJ10716 /FL=gb:NM_018191.1	NM_018191		NP_060661
218354_at	1.26E-04	gb:NM_016209.1 /DEF=Homo sapiens unknown (LOC51693), mRNA. /FEA=mRNA /GEN=LOC51693 /PROD=unknown /DB_XREF=gi:7706428 /UG=Hs.27445 unknown /FL=gb:AF089106.1 gb:AF161524.1 gb:NM_016209.1	NM_016209		NP_057293
218357_s_at	2.05E-04	gb:NM_012459.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 8 (yeast) homolog B (TIMM8B), mRNA. /FEA=mRNA /GEN=TIMM8B /PROD=translocase of inner mitochondrial membrane 8(yeast) homolog B /DB_XREF=gi:6912711 /UG=Hs.279915 translocase of inner mitochondrial membrane 8 (yeast) homolog B /FL=gb:AF150087.1 gb:AF152350.1 gb:NM_012459.1	NM_012459		NP_036591
218360_at	1.26E-04	gb:NM_020673.1 /DEF=Homo sapiens RAB22A, member RAS oncogene family (RAB22A), mRNA. /FEA=mRNA /GEN=RAB22A /PROD=RAB22A, member RAS oncogene family /DB_XREF=gi:10190713 /UG=Hs.288968 RAB22A, member RAS oncogene family /FL=gb:NM_020673.1 gb:AF091034.2	NM_020673		NP_065724

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218364_at	7.58E-04	gb:NM_017724.1 /DEF=Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2), mRNA. /FEA=mRNA /GEN=LRRFIP2 /PROD=leucine rich repeat (in FLII) interactingprotein 2 /DB_XREF=gi:8923223 /UG=Hs.57672 leucine rich repeat (in FLII) interacting protein 2 /FL=gb:NM_017724.1	NM_017724		NP_060194
218375_at	0.033533	gb:NM_024047.1 /DEF=Homo sapiens hypothetical protein MGC3037 (MGC3037), mRNA. /FEA=mRNA /GEN=MGC3037 /PROD=hypothetical protein MGC3037 /DB_XREF=gi:13129009 /UG=Hs.301789 hypothetical protein MGC3037 /FL=gb:BC000542.1 gb:NM_024047.1	NM_024047		NP_076952
218376_s_at	0.013223	gb:NM_022765.1 /DEF=Homo sapiens hypothetical protein FLJ11937 (FLJ11937), mRNA. /FEA=mRNA /GEN=FLJ11937 /PROD=hypothetical protein FLJ11937 /DB_XREF=gi:12232438 /UG=Hs.33476 hypothetical protein FLJ11937 /FL=gb:NM_022765.1	NM_022765		NP_073602
218379_at	0.007779	gb:NM_016090.1 /DEF=Homo sapiens RNA binding motif protein 7 (RBM7), mRNA. /FEA=mRNA /GEN=RBM7 /PROD=RNA binding motif protein 7 /DB_XREF=gi:9994184 /UG=Hs.5887 RNA binding motif protein 7 /FL=gb:AF156098.1 gb:NM_016090.1	NM_016090		NP_057174

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218383_at	0.001116	gb:NM_017815.1 /DEF=Homo sapiens hypothetical protein FLJ20424 (FLJ20424), mRNA. /FEA=mRNA /GEN=FLJ20424 /PROD=hypothetical protein FLJ20424 /DB_XREF=gi:8923395 /UG=Hs.8886 hypothetical protein FLJ20424 /FL=gb:BC002554.1 gb:BC001916.1 gb:NM_017815.1	NM_017815		NP_060285
218394_at	0.00161	gb:NM_024589.1 /DEF=Homo sapiens hypothetical protein FLJ22386 (FLJ22386), mRNA. /FEA=mRNA /GEN=FLJ22386 /PROD=hypothetical protein FLJ22386 /DB_XREF=gi:13375778 /UG=Hs.22795 hypothetical protein FLJ22386 /FL=gb:AL136675.1 gb:NM_024589.1	NM_024589		NP_078865
218399_s_at	0.033793	gb:NM_017955.1 /DEF=Homo sapiens hypothetical protein FLJ20764 (FLJ20764), mRNA. /FEA=mRNA /GEN=FLJ20764 /PROD=hypothetical protein FLJ20764 /DB_XREF=gi:8923673 /UG=Hs.34045 hypothetical protein FLJ20764 /FL=gb:NM_017955.1	NM_017955		NP_663747
218404_at	0.001116	gb:NM_013322.1 /DEF=Homo sapiens sorting nexin 10 (SNX10), mRNA. /FEA=mRNA /GEN=SNX10 /PROD=sorting nexin 10 /DB_XREF=gi:7019536 /UG=Hs.106260 sorting nexin 10 /FL=gb:AF121860.1 gb:NM_013322.1	NM_013322		NP_037454
218414_s_at	5.03E-04	gb:NM_017668.1 /DEF=Homo sapiens LIS1-interacting protein NUDE1, rat homolog (FLJ20101), mRNA. /FEA=mRNA /GEN=FLJ20101 /PROD=LIS1-interacting protein NUDE1, rat homolog /DB_XREF=gi:8923109 /UG=Hs.263925 LIS1-interacting protein NUDE1, rat homolog /FL=gb:BC001421.1 gb:NM_017668.1	NM_017668		NP_060138

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218419_s_at	0.02145	gb:NM_024107.1 /DEF=Homo sapiens hypothetical protein MGC3123 (MGC3123), mRNA. /FEA=mRNA /GEN=MGC3123 /PROD=hypothetical protein MGC3123 /DB_XREF=gi:13129117 /UG=Hs.288600 hypothetical protein MGC3123 /FL=gb:BC001792.1 gb:NM_024107.1	NM_024107		NP_803190
218420_s_at	0.037567	gb:NM_025138.1 /DEF=Homo sapiens hypothetical protein FLJ12661 (FLJ12661), mRNA. /FEA=mRNA /GEN=FLJ12661 /PROD=hypothetical protein FLJ12661 /DB_XREF=gi:13376720 /UG=Hs.318526 hypothetical protein FLJ12661 /FL=gb:NM_025138.1	NM_025138		NP_733837
218423_x_at	1.26E-04	gb:NM_016516.1 /DEF=Homo sapiens tumor antigen SLP-8p (HCC8), mRNA. /FEA=mRNA /GEN=HCC8 /PROD=tumor antigen SLP-8p /DB_XREF=gi:7705396 /UG=Hs.48499 tumor antigen SLP-8p /FL=gb:AF102177.1 gb:NM_016516.1	NM_016516		NP_057600
218424_s_at	0.008237	gb:NM_018234.1 /DEF=Homo sapiens hypothetical protein FLJ10829 (FLJ10829), mRNA. /FEA=mRNA /GEN=FLJ10829 /PROD=hypothetical protein FLJ10829 /DB_XREF=gi:8922696 /UG=Hs.57655 hypothetical protein FLJ10829 /FL=gb:NM_018234.1	NM_018234		NP_060704
218426_s_at	0.00161	gb:NM_019011.1 /DEF=Homo sapiens TRIAD3 protein (TRIAD3), mRNA. /FEA=mRNA /GEN=TRIAD3 /PROD=TRIAD3 protein /DB_XREF=gi:9507204 /UG=Hs.86228 TRIAD3 protein /FL=gb:BC000787.1 gb:NM_019011.1	NM_019011		NP_061884

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218431_at	0.02145	gb:NM_022067.1 /DEF=Homo sapiens hypothetical protein FLJ12707 (FLJ12707), mRNA. /FEA=mRNA /GEN=FLJ12707 /PROD=hypothetical protein FLJ12707 /DB_XREF=gi:11545778 /UG=Hs.16157 hypothetical protein FLJ12707 /FL=gb:NM_022067.1	NM_022067		NP_071350
218439_s_at	0.003817	gb:NM_016144.1 /DEF=Homo sapiens PTD002 protein (PTD002), mRNA. /FEA=mRNA /GEN=PTD002 /PROD=PTD002 protein /DB_XREF=gi:7706662 /UG=Hs.41767 PTD002 protein /FL=gb:BC005179.1 gb:AF078857.1 gb:NM_016144.1	NM_016144		NP_057228
218448_at	0.010205	gb:NM_017896.1 /DEF=Homo sapiens hypothetical protein FLJ20602 (FLJ20602), mRNA. /FEA=mRNA /GEN=FLJ20602 /PROD=hypothetical protein FLJ20602 /DB_XREF=gi:8923556 /UG=Hs.103808 hypothetical protein FLJ20602 /FL=gb:NM_017896.1	NM_017896		NP_060366
218450_at	0.004341	gb:NM_015987.1 /DEF=Homo sapiens heme-binding protein (HEBP), mRNA. /FEA=mRNA /GEN=HEBP /PROD=heme-binding protein /DB_XREF=gi:7705404 /UG=Hs.108675 heme-binding protein /FL=gb:AF117615.1 gb:NM_015987.1 gb:AF167473.1	NM_015987		NP_057071
218454_at	1.26E-04	gb:NM_024829.1 /DEF=Homo sapiens hypothetical protein FLJ22662 (FLJ22662), mRNA. /FEA=mRNA /GEN=FLJ22662 /PROD=hypothetical protein FLJ22662 /DB_XREF=gi:13376231 /UG=Hs.178470 hypothetical protein FLJ22662 /FL=gb:NM_024829.1	NM_024829		NP_079105

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218458_at	1.26E-04	gb:NM_022471.1 /DEF=Homo sapiens hypothetical protein FLJ13057 similar to germ cell-less (FLJ13057), mRNA. /FEA=mRNA /GEN=FLJ13057 /PROD=hypothetical protein FLJ13057 similar to germcell-less /DB_XREF=gi:11968020 /UG=Hs.243122 hypothetical protein FLJ13057 similar to germ cell-less /FL=gb:NM_022471.1	NM_022471		NP_848526
218465_at	2.05E-04	gb:NM_018126.1 /DEF=Homo sapiens hypothetical protein FLJ10525 (FLJ10525), mRNA. /FEA=mRNA /GEN=FLJ10525 /PROD=hypothetical protein FLJ10525 /DB_XREF=gi:8922490 /UG=Hs.31082 hypothetical protein FLJ10525 /FL=gb:BC000948.1 gb:NM_018126.1	NM_018126		NP_060596
218471_s_at	0.001886	gb:NM_024649.1 /DEF=Homo sapiens hypothetical protein FLJ23590 (FLJ23590), mRNA. /FEA=mRNA /GEN=FLJ23590 /PROD=hypothetical protein FLJ23590 /DB_XREF=gi:13375892 /UG=Hs.54890 hypothetical protein FLJ23590 /FL=gb:NM_024649.1	NM_024649		NP_078925
218472_s_at	5.03E-04	gb:NM_015946.1 /DEF=Homo sapiens pelota (Drosophila) homolog (PELO), mRNA. /FEA=mRNA /GEN=PELO /PROD=CGI-17 protein /DB_XREF=gi:7705599 /UG=Hs.5798 pelota (Drosophila) homolog /FL=gb:AF139828.1 gb:AF132951.1 gb:NM_015946.1	NM_015946		NP_057030
218482_at	2.05E-04	gb:NM_020189.1 /DEF=Homo sapiens DC6 protein (DC6), mRNA. /FEA=mRNA /GEN=DC6 /PROD=DC6 protein /DB_XREF=gi:9910185 /UG=Hs.283740 DC6 protein /FL=gb:AF201940.1 gb:AF173296.1 gb:NM_020189.1	NM_020189		NP_064574

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218487_at	2.05E-04	gb:BC000977.1 /DEF=Homo sapiens, aminolevulinate, delta-, dehydratase, clone MGC:5057, mRNA, complete cds. /FEA=mRNA /PROD=aminolevulinate, delta-, dehydratase /DB_XREF=gi:12654312 /UG=Hs.1227 aminolevulinate, delta-, dehydratase /FL=gb:BC000977.1 gb:M13928.1 gb:Nm_000031.1	BC000977		NP_000022
218490_s_at	0.022795	gb:Nm_018443.1 /DEF=Homo sapiens zinc finger protein ZNF140-like protein (LOC55828), mRNA. /FEA=mRNA /GEN=LOC55828 /PROD=zinc finger protein ZNF140-like protein /DB_XREF=gi:8923850 /UG=Hs.125287 zinc finger protein ZNF140-like protein /FL=gb:AF155656.1 gb:Nm_018443.1	NM_018443		NP_061145
218492_s_at	0.020793	gb:Nm_030573.1 /DEF=Homo sapiens hypothetical protein MGC10963 (MGC10963), mRNA. /FEA=mRNA /GEN=MGC10963 /PROD=hypothetical protein MGC10963 /DB_XREF=gi:13386483 /UG=Hs.14927 hypothetical protein MGC10963 /FL=gb:BC004346.1 gb:Nm_030573.1	NM_030573		NP_085050
218493_at	0.025958	gb:Nm_024571.1 /DEF=Homo sapiens hypothetical protein FLJ22940 (FLJ22940), mRNA. /FEA=mRNA /GEN=FLJ22940 /PROD=hypothetical protein FLJ22940 /DB_XREF=gi:13443017 /UG=Hs.15277 hypothetical protein FLJ22940 /FL=gb:BC001381.1 gb:Nm_024571.1	NM_024571		NP_078847

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218498_s_at	1.32E-04	gb:NM_014584.1 /DEF=Homo sapiens ERO1 (S. cerevisiae)-like (ERO1L), mRNA. /FEA=mRNA /GEN=ERO1L /PROD=ERO1 (S. cerevisiae)-like /DB_XREF=gi:7657068 /UG=Hs.25740 ERO1 (S. cerevisiae)-like /FL=gb:AF081886.1 gb:NM_014584.1	NM_014584		NP_055399
218499_at	0.026891	gb:NM_016542.1 /DEF=Homo sapiens serinethreonine protein kinase MASK (LOC51765), mRNA. /FEA=mRNA /GEN=LOC51765 /PROD=serinethreonine protein kinase MASK /DB_XREF=gi:7706568 /UG=Hs.23643 serinethreonine protein kinase MASK /FL=gb:AB040057.1 gb:NM_016542.1	NM_016542		NP_057626
218502_s_at	0.005852	gb:NM_014112.1 /DEF=Homo sapiens trichorhinophalangeal syndrome I gene (TRPS1), mRNA. /FEA=mRNA /GEN=TRPS1 /PROD=trichorhinophalangeal syndrome I gene /DB_XREF=gi:7657658 /UG=Hs.26102 trichorhinophalangeal syndrome I /FL=gb:AF183810.1 gb:NM_014112.1	NM_014112		NP_054831
218506_x_at	2.05E-04	gb:NM_018459.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM045 (BM045), mRNA. /FEA=mRNA /GEN=BM045 /PROD=uncharacterized bone marrow protein BM045 /DB_XREF=gi:8922103 /UG=Hs.8750 uncharacterized bone marrow protein BM045 /FL=gb:AF217521.1 gb:NM_018459.1	NM_018459		NP_060929

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218508_at	5.64E-04	gb:NM_018403.1 /DEF=Homo sapiens transcription factor (SMIF gene) (HSA275986), mRNA. /FEA=mRNA /GEN=HSA275986 /PROD=transcription factor (SMIF gene) /DB_XREF=gi:8923766 /UG=Hs.71414 transcription factor (SMIF gene) /FL=gb:NM_018403.1	NM_018403		NP_060873
218514_at	3.26E-04	gb:NM_018149.1 /DEF=Homo sapiens hypothetical protein FLJ10587 (FLJ10587), mRNA. /FEA=mRNA /GEN=FLJ10587 /PROD=hypothetical protein FLJ10587 /DB_XREF=gi:8922539 /UG=Hs.7296 hypothetical protein FLJ10587 /FL=gb:NM_018149.1	NM_018149		NP_060619
218518_at	0.016934	gb:NM_016603.1 /DEF=Homo sapiens GAP-like protein (LOC51306), mRNA. /FEA=mRNA /GEN=LOC51306 /PROD=GAP-like protein /DB_XREF=gi:7706136 /UG=Hs.82035 potential nuclear protein C5ORF5; GAP-like protein /FL=gb:AF251038.1 gb:AF157316.1 gb:NM_016603.1	NM_016603		NP_057687
218519_at	0.005852	gb:NM_017945.1 /DEF=Homo sapiens hypothetical protein FLJ20730 (FLJ20730), mRNA. /FEA=mRNA /GEN=FLJ20730 /PROD=hypothetical protein FLJ20730 /DB_XREF=gi:8923656 /UG=Hs.237480 hypothetical protein FLJ20730 /FL=gb:BC005207.1 gb:NM_017945.1	NM_017945		NP_060415
218520_at	0.005852	gb:NM_013254.1 /DEF=Homo sapiens TANK-binding kinase 1 (TBK1), mRNA. /FEA=mRNA /GEN=TBK1 /PROD=TANK-binding kinase 1 /DB_XREF=gi:7019546 /UG=Hs.21712 TANK-binding kinase 1 /FL=gb:AF191838.1 gb:NM_013254.1 gb:AF174536.1	NM_013254		NP_037386

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218522_s_at	0.001116	gb:NM_018174.1 /DEF=Homo sapiens hypothetical protein FLJ10669 (FLJ10669), mRNA. /FEA=mRNA /GEN=FLJ10669 /PROD=hypothetical protein FLJ10669 /DB_XREF=gi:8922582 /UG=Hs.66048 hypothetical protein FLJ10669 /FL=gb:NM_018174.1	NM_018174		NP_060644
218526_s_at	0.029453	gb:NM_014185.1 /DEF=Homo sapiens HSPC165 protein (HSPC165), mRNA. /FEA=mRNA /GEN=HSPC165 /PROD=HSPC165 protein /DB_XREF=gi:7661825 /UG=Hs.13605, HSPC165 protein /FL=gb:AF161514.1 gb:AF151070.1 gb:NM_014185.1 gb:NM_016492.1 gb:AF168714.1 gb:AF265206.1	NM_014185		NP_057576
218528_s_at	1.26E-04	gb:NM_022781.1 /DEF=Homo sapiens hypothetical protein FLJ21343 (FLJ21343), mRNA. /FEA=mRNA /GEN=FLJ21343 /PROD=hypothetical protein FLJ21343 /DB_XREF=gi:12232470 /UG=Hs.77823 hypothetical protein FLJ21343 /FL=gb:AL136817.1 gb:NM_022781.1	NM_022781		NP_073618
218531_at	0.013223	gb:NM_025124.1 /DEF=Homo sapiens hypothetical protein FLJ21749 (FLJ21749), mRNA. /FEA=mRNA /GEN=FLJ21749 /PROD=hypothetical protein FLJ21749 /DB_XREF=gi:13376700 /UG=Hs.288761 hypothetical protein FLJ21749 /FL=gb:NM_025124.1	NM_025124		NP_079400
218534_s_at	7.58E-04	gb:NM_018046.1 /DEF=Homo sapiens hypothetical protein FLJ10283 (FLJ10283), mRNA. /FEA=mRNA /GEN=FLJ10283 /PROD=hypothetical protein FLJ10283 /DB_XREF=gi:8922325 /UG=Hs.284216 hypothetical protein FLJ10283 /FL=gb:NM_018046.1	NM_018046		NP_060516

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218539_at	0.026891	gb:NM_017943.1 /DEF=Homo sapiens hypothetical protein FLJ20725 (FLJ20725), mRNA. /FEA=mRNA /GEN=FLJ20725 /PROD=hypothetical protein FLJ20725 /DB_XREF=gi:8923650 /UG=Hs.15467 hypothetical protein FLJ20725 /FL=gb:NM_017943.1	NM_017943		NP_060413
218549_s_at	1.26E-04	gb:NM_016033.1 /DEF=Homo sapiens CGI-90 protein (LOC51115), mRNA. /FEA=mRNA /GEN=LOC51115 /PROD=CGI-90 protein /DB_XREF=gi:7705802 /UG=Hs.44222 CGI-90 protein /FL=gb:AF151848.1 gb:NM_016033.1	NM_016033		NP_057117
218556_at	1.26E-04	gb:NM_014182.1 /DEF=Homo sapiens HSPC160 protein (HSPC160), mRNA. /FEA=mRNA /GEN=HSPC160 /PROD=HSPC160 protein /DB_XREF=gi:7661819 /UG=Hs.13144 HSPC160 protein /FL=gb:AF161509.1 gb:NM_014182.1	NM_014182		NP_054901
218563_at	0.013223	gb:NM_004542.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) (NDUFA3), mRNA. /FEA=mRNA /GEN=NDUFA3 /PROD=NADH dehydrogenase (ubiquinone) 1 alphasubcomplex, 3 (9kD, B9) /DB_XREF=gi:4758771 /UG=Hs.198269 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) /FL=gb:AF044955.1 gb:AF070653.1 gb:NM_004542.1	NM_004542		NP_004533
218572_at	0.021435	gb:NM_014169.1 /DEF=Homo sapiens HSPC134 protein (HSPC134), mRNA. /FEA=mRNA /GEN=HSPC134 /PROD=HSPC134 protein /DB_XREF=gi:7661793 /UG=Hs.279761 HSPC134 protein /FL=gb:AF212243.1 gb:AF161483.1 gb:NM_014169.1	NM_014169		NP_054888

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218578_at	1.32E-04	gb:NM_024529.1 /DEF=Homo sapiens hypothetical protein FLJ23316 (FLJ23316), mRNA. /FEA=mRNA /GEN=FLJ23316 /PROD=hypothetical protein FLJ23316 /DB_XREF=gi:13375677 /UG=Hs.5722 hypothetical protein FLJ23316 /FL=gb:AF312865.1 gb:NM_024529.1	NM_024529		NP_078805
218580_x_at	0.007779	gb:NM_017900.1 /DEF=Homo sapiens hypothetical protein FLJ20608 (FLJ20608), mRNA. /FEA=mRNA /GEN=FLJ20608 /PROD=hypothetical protein FLJ20608 /DB_XREF=gi:8923564 /UG=Hs.76239 hypothetical protein FLJ20608 /FL=gb:NM_017900.1	NM_017900		NP_060370
218582_at	2.05E-04	gb:NM_017824.1 /DEF=Homo sapiens hypothetical protein FLJ20445 (FLJ20445), mRNA. /FEA=mRNA /GEN=FLJ20445 /PROD=hypothetical protein FLJ20445 /DB_XREF=gi:8923414 /UG=Hs.10340 hypothetical protein FLJ20445 /FL=gb:NM_017824.1	NM_017824		NP_060294
218583_s_at	2.05E-04	gb:NM_020640.1 /DEF=Homo sapiens RP42 homolog (RP42), mRNA. /FEA=mRNA /GEN=RP42 /PROD=RP42 homolog /DB_XREF=gi:10190677 /UG=Hs.104613 RP42 homolog /FL=gb:NM_020640.1 gb:AF292100.2	NM_020640		NP_065691
218586_at	0.010205	gb:NM_018270.1 /DEF=Homo sapiens hypothetical protein FLJ10914 (FLJ10914), mRNA. /FEA=mRNA /GEN=FLJ10914 /PROD=hypothetical protein FLJ10914 /DB_XREF=gi:8922763 /UG=Hs.143954 hypothetical protein FLJ10914 /FL=gb:NM_018270.1	NM_018270		NP_060740

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218588_s_at	0.00161	gb:NM_018691.1 /DEF=Homo sapiens chromosome 5 open reading frame 3 (C5ORF3), mRNA. /FEA=mRNA /GEN=C5ORF3 /PROD=hypothetical protein /DB_XREF=gi:8922068 /UG=Hs.166551 chromosome 5 open reading frame 3 /FL=gb:NM_018691.1	NM_018691		NP_061161
218597_s_at	0.001477	gb:NM_018464.1 /DEF=Homo sapiens uncharacterized hematopoietic stemprogenitor cells protein MDS029 (MDS029), mRNA. /FEA=mRNA /GEN=MDS029 /PROD=uncharacterized hematopoietic stemprogenitorcells protein MDS029 /DB_XREF=gi:8923929 /UG=Hs.43549 uncharacterized hematopoietic stemprogenitor cells protein MDS029 /FL=gb:AF220049.1 gb:NM_018464.1	NM_018464		NP_060934
218603_at	0.001116	gb:NM_016217.1 /DEF=Homo sapiens hHDC for homolog of Drosophila headcase (LOC51696), mRNA. /FEA=mRNA /GEN=LOC51696 /PROD=hHDC for homolog of Drosophila headcase /DB_XREF=gi:7706434 /UG=Hs.6679 hHDC for homolog of Drosophila headcase /FL=gb:AB033492.1 gb:NM_016217.1	NM_016217		NP_057301
218606_at	1.26E-04	gb:NM_017740.1 /DEF=Homo sapiens hypothetical protein FLJ20279 (FLJ20279), mRNA. /FEA=mRNA /GEN=FLJ20279 /PROD=hypothetical protein FLJ20279 /DB_XREF=gi:8923254 /UG=Hs.9725 hypothetical protein FLJ20279 /FL=gb:NM_017740.1	NM_017740		NP_060210

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218616_at	2.05E-04	gb:NM_020395.1 /DEF=Homo sapiens hypothetical nuclear factor SBBI22 (LOC57117), mRNA. /FEA=mRNA /GEN=LOC57117 /PROD=hypothetical nuclear factor SBBI22 /DB_XREF=gi:9966872 /UG=Hs.26323 hypothetical nuclear factor SBBI22 /FL=gb:AF242524.1 gb:NM_020395.1	NM_020395		NP_065128
218617_at	0.035611	gb:NM_017646.1 /DEF=Homo sapiens tRNA isopentenylpyrophosphate transferase (IPT), mRNA. /FEA=mRNA /GEN=IPT /PROD=tRNA isopentenylpyrophosphate transferase /DB_XREF=gi:8923064 /UG=Hs.288036 tRNA isopentenylpyrophosphate transferase /FL=gb:NM_017646.1	NM_017646		NP_060116
218625_at	0.002864	gb:NM_016588.1 /DEF=Homo sapiens neuritin (LOC51299), mRNA. /FEA=mRNA /GEN=LOC51299 /PROD=neuritin /DB_XREF=gi:7706122 /UG=Hs.103291 neuritin /FL=gb:BC002683.1 gb:AF136631.1 gb:NM_016588.1	NM_016588		NP_057672
218627_at	1.26E-04	gb:NM_018370.1 /DEF=Homo sapiens hypothetical protein FLJ11259 (FLJ11259), mRNA. /FEA=mRNA /GEN=FLJ11259 /PROD=hypothetical protein FLJ11259 /DB_XREF=gi:8922957 /UG=Hs.184465 hypothetical protein FLJ11259 /FL=gb:NM_018370.1	NM_018370		NP_060840

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218628_at	0.001116	gb:NM_016053.1 /DEF=Homo sapiens CGI-116 protein (LOC51019), mRNA. /FEA=mRNA /GEN=LOC51019 /PROD=CGI-116 protein /DB_XREF=gi:7705621 /UG=Hs.18885 CGI-116 protein /FL=gb:AF151874.1 gb:NM_016053.1	NM_016053		NP_057137
218632_at	0.026891	gb:NM_024602.1 /DEF=Homo sapiens hypothetical protein FLJ21156 (FLJ21156), mRNA. /FEA=mRNA /GEN=FLJ21156 /PROD=hypothetical protein FLJ21156 /DB_XREF=gi:13375804 /UG=Hs.26058 hypothetical protein FLJ21156 /FL=gb:NM_024602.1	NM_024602		NP_078878
218637_at	0.033381	gb:NM_018439.1 /DEF=Homo sapiens hypothetical protein IMPACT (IMPACT), mRNA. /FEA=mRNA /GEN=IMPACT /PROD=hypothetical protein IMPACT /DB_XREF=gi:8923818 /UG=Hs.284245 hypothetical protein IMPACT /FL=gb:AF208694.1 gb:AB026264.1 gb:NM_018439.1	NM_018439		NP_060909
218639_s_at	0.035171	gb:NM_025112.1 /DEF=Homo sapiens hypothetical protein MGC11349 (MGC11349), mRNA. /FEA=mRNA /GEN=MGC11349 /PROD=hypothetical protein MGC11349 /DB_XREF=gi:13376684 /UG=Hs.288697 hypothetical protein MGC11349 /FL=gb:BC002940.1 gb:NM_025112.1	NM_025112		NP_079388

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218645_at	0.046906	gb:NM_021994.1 /DEF=Homo sapiens zinc finger protein 277 (ZNF277), mRNA. /FEA=mRNA /GEN=ZNF277 /PROD=zinc finger protein 277 /DB_XREF=gi:11496268 /UG=Hs.42636 zinc finger protein 277 /FL=gb:NM_021994.1 gb:AF209198.1	NM_021994		NP_068834
218652_s_at	0.007779	gb:NM_017733.1 /DEF=Homo sapiens hypothetical protein FLJ20265 (FLJ20265), mRNA. /FEA=mRNA /GEN=FLJ20265 /PROD=hypothetical protein FLJ20265 /DB_XREF=gi:8923239 /UG=Hs.7099 hypothetical protein FLJ20265 /FL=gb:BC001249.1 gb:BC000937.2 gb:NM_017733.1	NM_017733		NP_060203
218660_at	3.26E-04	gb:NM_003494.1 /DEF=Homo sapiens dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive) (DYSF), mRNA. /FEA=mRNA /GEN=DYSF /PROD=dysferlin /DB_XREF=gi:4503430 /UG=Hs.143897 dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive) /FL=gb:AF075575.1 gb:NM_003494.1	NM_003494		NP_003485
218661_at	5.64E-04	gb:NM_024845.1 /DEF=Homo sapiens hypothetical protein FLJ14154 (FLJ14154), mRNA. /FEA=mRNA /GEN=FLJ14154 /PROD=hypothetical protein FLJ14154 /DB_XREF=gi:13376260 /UG=Hs.184870 hypothetical protein FLJ14154 /FL=gb:NM_024845.1	NM_024845		NP_079121

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218662_s_at	2.05E-04	gb:NM_022346.1 /DEF=Homo sapiens chromosome condensation protein G (HCAP-G), mRNA. /FEA=mRNA /GEN=HCAP-G /PROD=chromosome condensation protein G /DB_XREF=gi:11641252 /UG=Hs.193602 chromosome condensation protein G /FL=gb:AF235023.1 gb:NM_022346.1 gb:AF331796.1 gb:BC000827.1 gb:AB013299.1	NM_022346		NP_071741
218668_s_at	3.26E-04	gb:NM_021183.1 /DEF=Homo sapiens hypothetical protein similar to small G proteins, especially RAP-2A (LOC57826), mRNA. /FEA=mRNA /GEN=LOC57826 /PROD=hypothetical protein similar to small Gproteins, especially RAP-2A /DB_XREF=gi:10880976 /UG=Hs.225979 hypothetical protein similar to small G proteins, especially RAP-2A /FL=gb:NM_021183.1 gb:BC003403.1	NM_021183		NP_067006
218669_at	0.00161	gb:NM_021183.1 /DEF=Homo sapiens hypothetical protein similar to small G proteins, especially RAP-2A (LOC57826), mRNA. /FEA=mRNA /GEN=LOC57826 /PROD=hypothetical protein similar to small Gproteins, especially RAP-2A /DB_XREF=gi:10880976 /UG=Hs.225979 hypothetical protein similar to small G proteins, especially RAP-2A /FL=gb:NM_021183.1 gb:BC003403.1	NM_021183		NP_067006

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218673_s_at	0.041055	gb:NM_006395.1 /DEF=Homo sapiens ubiquitin activating enzyme E1-like protein (GSA7), mRNA. /FEA=mRNA /GEN=GSA7 /PROD=ubiquitin activating enzyme E1-like protein /DB_XREF=gi:5453667 /UG=Hs.278607 ubiquitin activating enzyme E1-like protein /FL=gb:AF094516.1 gb:NM_006395.1	NM_006395		NP_006386
218676_s_at	0.005852	gb:NM_021213.1 /DEF=Homo sapiens phosphatidylcholine transfer protein (PCTP), mRNA. /FEA=mRNA /GEN=PCTP /PROD=phosphatidylcholine transfer protein /DB_XREF=gi:10864026 /UG=Hs.285218 phosphatidylcholine transfer protein /FL=gb:NM_021213.1 gb:AF151638.1	NM_021213		NP_067036
218679_s_at	0.004341	gb:NM_016208.1 /DEF=Homo sapiens VPS28 protein (LOC51160), mRNA. /FEA=mRNA /GEN=LOC51160 /PROD=VPS28 protein /DB_XREF=gi:7705884 /UG=Hs.293441 VPS28 protein /FL=gb:AF316887.1 gb:AF182844.1 gb:NM_016208.1	NM_016208		NP_057292
218682_s_at	1.26E-04	gb:NM_018158.1 /DEF=Homo sapiens hypothetical protein FLJ10624 (FLJ10624), mRNA. /FEA=mRNA /GEN=FLJ10624 /PROD=hypothetical protein FLJ10624 /DB_XREF=gi:8922556 /UG=Hs.306000 hypothetical protein FLJ10624 /FL=gb:NM_018158.1	NM_018158		NP_060628
218698_at	0.02145	gb:NM_015957.1 /DEF=Homo sapiens CGI-29 protein (LOC51074), mRNA. /FEA=mRNA /GEN=LOC51074 /PROD=CGI-29 protein /DB_XREF=gi:7705723 /UG=Hs.104058 CGI-29 protein /FL=gb:AF132963.1 gb:NM_015957.1	NM_015957		NP_057041

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218703_at	0.021448	gb:NM_012430.1 /DEF=Homo sapiens sec22 homolog (SEC22A), mRNA. /FEA=mRNA /GEN=SEC22A /PROD=sec22 homolog /DB_XREF=gi:6912647 /UG=Hs.183655 sec22 homolog /FL=gb:AF100749.1 gb:NM_012430.1	NM_012430		NP_036562
218718_at	3.26E-04	gb:NM_016205.1 /DEF=Homo sapiens platelet derived growth factor C (PDGFC), mRNA. /FEA=mRNA /GEN=PDGFC /PROD=secretory growth factor-like protein fallotein /DB_XREF=gi:9994186 /UG=Hs.43080 platelet derived growth factor C /FL=gb:AF091434.1 gb:AF244813.1 gb:AB033831.1 gb:NM_016205.1	NM_016205		NP_057289
218722_s_at	0.025245	gb:NM_024661.1 /DEF=Homo sapiens hypothetical protein FLJ12436 (FLJ12436), mRNA. /FEA=mRNA /GEN=FLJ12436 /PROD=hypothetical protein FLJ12436 /DB_XREF=gi:13375914 /UG=Hs.69485 hypothetical protein FLJ12436 /FL=gb:NM_024661.1	NM_024661		NP_078937
218723_s_at	0.016934	gb:NM_014059.1 /DEF=Homo sapiens RGC32 protein (RGC32), mRNA. /FEA=mRNA /GEN=RGC32 /PROD=RGC32 protein /DB_XREF=gi:7662650 /UG=Hs.76640 RGC32 protein /FL=gb:AF036549.1 gb:NM_014059.1	NM_014059		NP_054778
218724_s_at	0.003539	gb:NM_021809.1 /DEF=Homo sapiens TGF(beta)-induced transcription factor 2 (TGIF2), mRNA. /FEA=mRNA /GEN=TGIF2 /PROD=TGF(beta)-induced transcription factor 2 /DB_XREF=gi:11141880 /UG=Hs.94785 TGF(beta)-induced transcription factor 2 /FL=gb:AB042646.1 gb:NM_021809.1	NM_021809		NP_068581

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218728_s_at	1.26E-04	gb:NM_014184.1 /DEF=Homo sapiens HSPC163 protein (HSPC163), mRNA. /FEA=mRNA /GEN=HSPC163 /PROD=HSPC163 protein /DB_XREF=gi:7661823 /UG=Hs.108854 HSPC163 protein /FL=gb:BC000573.1 gb:AF161512.1 gb:NM_014184.1	NM_014184		NP_054903
218729_at	0.007258	gb:NM_020169.1 /DEF=Homo sapiens latexin protein (LXN), mRNA. /FEA=mRNA /GEN=LXN /PROD=latexin protein /DB_XREF=gi:9910395 /UG=Hs.109276 latexin protein /FL=gb:BC005346.1 gb:AF282626.1 gb:NM_020169.1	NM_020169		NP_064554
218734_at	0.004341	gb:NM_024771.1 /DEF=Homo sapiens hypothetical protein FLJ13848 (FLJ13848), mRNA. /FEA=mRNA /GEN=FLJ13848 /PROD=hypothetical protein FLJ13848 /DB_XREF=gi:13376118 /UG=Hs.136976 hypothetical protein FLJ13848 /FL=gb:NM_024771.1	NM_024771		NP_079047
218738_s_at	5.03E-04	gb:NM_016271.1 /DEF=Homo sapiens STRIN protein (STRIN), mRNA. /FEA=mRNA /GEN=STRIN /PROD=STRIN protein /DB_XREF=gi:7706722 /UG=Hs.180403 STRIN protein /FL=gb:AF162680.3 gb:NM_016271.1	NM_016271		NP_057355
218739_at	2.05E-04	gb:NM_016006.1 /DEF=Homo sapiens CGI-58 protein (LOC51099), mRNA. /FEA=mRNA /GEN=LOC51099 /PROD=CGI-58 protein /DB_XREF=gi:7705770 /UG=Hs.19385 CGI-58 protein /FL=gb:AF151816.1 gb:NM_016006.1	NM_016006		NP_057090

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218761_at	7.58E-04	gb:NM_017610.1 /DEF=Homo sapiens hypothetical protein DKFZp761D081 (DKFZp761D081), mRNA. /FEA=mRNA /GEN=DKFZp761D081 /PROD=hypothetical protein DKFZp761D081 /DB_XREF=gi:8922164 /UG=Hs.12504 hypothetical protein DKFZp761D081 /FL=gb:NM_017610.1	NM_017610		NP_060080
218764_at	5.03E-04	gb:NM_024064.1 /DEF=Homo sapiens hypothetical protein MGC5363 (MGC5363), mRNA. /FEA=mRNA /GEN=MGC5363 /PROD=hypothetical protein MGC5363 /DB_XREF=gi:13129041 /UG=Hs.1880 hypothetical protein MGC5363 /FL=gb:BC001000.2 gb:NM_024064.1	NM_024064		
218773_s_at	7.58E-04	gb:NM_012228.1 /DEF=Homo sapiens pilin-like transcription factor (PILB), mRNA. /FEA=mRNA /GEN=PILB /PROD=pilin-like transcription factor /DB_XREF=gi:6912591 /UG=Hs.279754 pilin-like transcription factor /FL=gb:AF122004.1 gb:NM_012228.1	NM_012228		NP_036360
218782_s_at	0.013223	gb:NM_014109.1 /DEF=Homo sapiens PRO2000 protein (PRO2000), mRNA. /FEA=mRNA /GEN=PRO2000 /PROD=PRO2000 protein /DB_XREF=gi:7662630 /UG=Hs.46677 PRO2000 protein /FL=gb:AF118088.1 gb:NM_014109.1	NM_014109		NP_054828
218789_s_at	0.001848	gb:NM_019021.1 /DEF=Homo sapiens hypothetical protein (FLJ20010), mRNA. /FEA=mRNA /GEN=FLJ20010 /PROD=hypothetical protein /DB_XREF=gi:9506646 /UG=Hs.91816 hypothetical protein /FL=gb:NM_019021.1	NM_019021		NP_061894

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218799_at	0.001205	gb:NM_018066.1 /DEF=Homo sapiens hypothetical protein FLJ10349 (FLJ10349), mRNA. /FEA=mRNA /GEN=FLJ10349 /PROD=hypothetical protein FLJ10349 /DB_XREF=gi:8922365 /UG=Hs.19333 hypothetical protein FLJ10349 /FL=gb:NM_018066.1	NM_018066		NP_060536
218805_at	0.001116	gb:NM_018384.1 /DEF=Homo sapiens hypothetical protein FLJ11296 (FLJ11296), mRNA. /FEA=mRNA /GEN=FLJ11296 /PROD=hypothetical protein FLJ11296 /DB_XREF=gi:8922984 /UG=Hs.26194 hypothetical protein FLJ11296 /FL=gb:NM_018384.1	NM_018384		NP_060854
218806_s_at	2.32E-04	gb:AF118887.1 /DEF=Homo sapiens VAV-3 protein (VAV-3) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=VAV-3 /PROD=VAV-3 protein /DB_XREF=gi:4416407 /UG=Hs.267659 vav 3 oncogene /FL=gb:AF067817.1 gb:AF118887.1 gb:NM_006113.2	AF118887		NP_006104
218809_at	0.005852	gb:NM_024960.1 /DEF=Homo sapiens hypothetical protein FLJ11729 (FLJ11729), mRNA. /FEA=mRNA /GEN=FLJ11729 /PROD=hypothetical protein FLJ11729 /DB_XREF=gi:13376450 /UG=Hs.286212 hypothetical protein FLJ11729 /FL=gb:NM_024960.1	NM_024960		NP_705905
218827_s_at	0.016074	gb:NM_018069.1 /DEF=Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA. /FEA=mRNA /GEN=FLJ10352 /PROD=hypothetical protein FLJ10352 /DB_XREF=gi:8922371 /UG=Hs.100914 hypothetical protein FLJ10352 /FL=gb:NM_018069.1	NM_018069		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218829_s_at	1.26E-04	gb:NM_017780.1 /DEF=Homo sapiens hypothetical protein FLJ20357 (FLJ20357), mRNA. /FEA=mRNA /GEN=FLJ20357 /PROD=hypothetical protein FLJ20357 /DB_XREF=gi:8923329 /UG=Hs.105461 hypothetical protein FLJ20357 /FL=gb:NM_017780.1	NM_017780		
218833_at	3.09E-04	gb:NM_016653.1 /DEF=Homo sapiens sterile-alpha motif and leucine zipper containing kinase AZK (ZAK), mRNA. /FEA=mRNA /GEN=ZAK /PROD=sterile-alpha motif and leucine zippercontaining kinase AZK /DB_XREF=gi:7706600 /UG=Hs.115175 sterile-alpha motif and leucine zipper containing kinase AZK /FL=gb:AB049733.1 gb:AF238255.1 gb:AF251441.1 gb:NM_016653.1 gb:AB030034.1	NM_016653		NP_598407
218841_at	0.00161	gb:NM_024095.1 /DEF=Homo sapiens hypothetical protein MGC5540 (MGC5540), mRNA. /FEA=mRNA /GEN=MGC5540 /PROD=hypothetical protein MGC5540 /DB_XREF=gi:13129097 /UG=Hs.267400 hypothetical protein MGC5540 /FL=gb:BC001321.1 gb:NM_024095.1	NM_024095		NP_077000
218846_at	7.58E-04	gb:NM_004830.1 /DEF=Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 3 (130kD) (CRSP3), mRNA. /FEA=mRNA /GEN=CRSP3 /PROD=cofactor required for Sp1 transcriptionalactivation, subunit 3 (130kD) /DB_XREF=gi:7019352 /UG=Hs.29679 cofactor required for Sp1 transcriptional activation, subunit 3 (130kD) /FL=gb:AF135022.1 gb:AF105332.1 gb:NM_004830.1	NM_004830		NP_057063

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218850_s_at	0.020793	gb:NM_014240.1 /DEF=Homo sapiens LIM domains containing 1 (LIMD1), mRNA. /FEA=mRNA /GEN=LIMD1 /PROD=LIM domains containing 1 /DB_XREF=gi:7657306 /UG=Hs.48469 LIM domains containing 1 /FL=gb:NM_014240.1	NM_014240		NP_055055
218852_at	0.003171	gb:NM_017917.1 /DEF=Homo sapiens hypothetical protein FLJ20644 (FLJ20644), mRNA. /FEA=mRNA /GEN=FLJ20644 /PROD=hypothetical protein FLJ20644 /DB_XREF=gi:8923599 /UG=Hs.49376 hypothetical protein FLJ20644 /FL=gb:NM_017917.1	NM_017917		NP_060387
218853_s_at	0.004341	gb:NM_019556.1 /DEF=Homo sapiens hypothetical protein dJ473B4 (DJ473B4), mRNA. /FEA=mRNA /GEN=DJ473B4 /PROD=hypothetical protein dJ473B4 /DB_XREF=gi:9506542 /UG=Hs.57549 hypothetical protein dJ473B4 /FL=gb:NM_019556.1	NM_019556		NP_062456
218854_at	5.03E-04	gb:NM_013352.1 /DEF=Homo sapiens squamous cell carcinoma antigen recognized by T cell (SART-2), mRNA. /FEA=mRNA /GEN=SART-2 /PROD=squamous cell carcinoma antigen recognized by T cell /DB_XREF=gi:7019520 /UG=Hs.58636 squamous cell carcinoma antigen recognized by T cell /FL=gb:AF098066.1 gb:NM_013352.1	NM_013352		NP_037484

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218870_at	0.026891	gb:NM_018460.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM046 (BM046), mRNA. /FEA=mRNA /GEN=BM046 /PROD=uncharacterized bone marrow protein BM046 /DB_XREF=gi:8922105 /UG=Hs.177812 uncharacterized bone marrow protein BM046 /FL=gb:AF217507.1 gb:AF217522.1 gb:NM_018460.1 gb:AF212222.1	NM_018460		NP_060930
218871_x_at	2.05E-04	gb:NM_018590.1 /DEF=Homo sapiens hypothetical protein PRO0082 (PRO0082), mRNA. /FEA=mRNA /GEN=PRO0082 /PROD=hypothetical protein PRO0082 /DB_XREF=gi:8923965 /UG=Hs.180758 hypothetical protein PRO0082 /FL=gb:AF116646.1 gb:NM_018590.1	NM_018590		NP_061060
218872_at	2.75E-04	gb:NM_017899.1 /DEF=Homo sapiens hypothetical protein FLJ20607 (FLJ20607), mRNA. /FEA=mRNA /GEN=FLJ20607 /PROD=hypothetical protein FLJ20607 /DB_XREF=gi:8923562 /UG=Hs.18791 hypothetical protein FLJ20607 /FL=gb:NM_017899.1	NM_017899		NP_060369
218873_at	0.041055	gb:NM_017710.1 /DEF=Homo sapiens hypothetical protein FLJ20203 (FLJ20203), mRNA. /FEA=mRNA /GEN=FLJ20203 /PROD=hypothetical protein FLJ20203 /DB_XREF=gi:8923193 /UG=Hs.20594 hypothetical protein FLJ20203 /FL=gb:NM_017710.1	NM_017710		NP_115668
218880_at	0.010205	FOS-like antigen 2	N36408	Hs.301612	NP_005244

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218882_s_at	0.041055	gb:NM_006784.1 /DEF=Homo sapiens WD repeat domain 3 (WDR3), mRNA. /FEA=mRNA /GEN=WDR3 /PROD=WD repeat-containing protein 3 /DB_XREF=gi:5803220 /UG=Hs.33085 WD repeat domain 3 /FL=gb:AF083217.1 gb:NM_006784.1	NM_006784		NP_006775
218883_s_at	2.05E-04	gb:NM_024629.1 /DEF=Homo sapiens hypothetical protein FLJ23468 (FLJ23468), mRNA. /FEA=mRNA /GEN=FLJ23468 /PROD=hypothetical protein FLJ23468 /DB_XREF=gi:13375855 /UG=Hs.38178 hypothetical protein FLJ23468 /FL=gb:NM_024629.1	NM_024629		NP_078905
218888_s_at	0.048086	gb:NM_018092.1 /DEF=Homo sapiens hypothetical protein FLJ10430 (FLJ10430), mRNA. /FEA=mRNA /GEN=FLJ10430 /PROD=hypothetical protein FLJ10430 /DB_XREF=gi:8922419 /UG=Hs.6823 hypothetical protein FLJ10430 /FL=gb:NM_018092.1	NM_018092		NP_060562
218896_s_at	0.001116	gb:NM_018553.1 /DEF=Homo sapiens ELG protein (HSA277841), mRNA. /FEA=mRNA /GEN=HSA277841 /PROD=ELG protein /DB_XREF=gi:8923770 /UG=Hs.120963 ELG protein /FL=gb:NM_018553.1	NM_018553		NP_061023
218904_s_at	1.92E-04	gb:NM_017998.1 /DEF=Homo sapiens hypothetical protein FLJ10110 (FLJ10110), mRNA. /FEA=mRNA /GEN=FLJ10110 /PROD=hypothetical protein FLJ10110 /DB_XREF=gi:8922232 /UG=Hs.264363 hypothetical protein FLJ10110 /FL=gb:NM_017998.1	NM_017998		NP_060468

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218909_at	0.027243	gb:NM_012424.1 /DEF=Homo sapiens ribosomal protein S6 kinase, 52kD, polypeptide 1 (RPS6KC1), mRNA. /FEA=mRNA /GEN=RPS6KC1 /PROD=ribosomal protein S6 kinase, 52kD, polypeptide1 /DB_XREF=gi:6912635 /UG=Hs.30352 ribosomal protein S6 kinase, 52kD, polypeptide 1 /FL=gb:AF037447.1 gb:NM_012424.1	NM_012424		NP_036556
218917_s_at	1.26E-04	gb:NM_018450.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM029 (BM029), mRNA. /FEA=mRNA /GEN=BM029 /PROD=uncharacterized bone marrow protein BM029 /DB_XREF=gi:8922086 /UG=Hs.8740 uncharacterized bone marrow protein BM029 /FL=gb:AF217506.1 gb:NM_018450.1	NM_018450		NP_624361
218921_at	0.022795	gb:NM_021805.1 /DEF=Homo sapiens single Ig IL-1R-related molecule (SIGIRR), mRNA. /FEA=mRNA /GEN=SIGIRR /PROD=single Ig IL-1R-related molecule /DB_XREF=gi:11141876 /UG=Hs.11809 single Ig IL-1R-related molecule /FL=gb:NM_021805.1 gb:BC003591.1	NM_021805		NP_068577
218923_at	0.001116	chitobiase, di-N-acetyl-	AW304174	Hs.135578	NP_004379
218930_s_at	0.00161	gb:NM_018374.1 /DEF=Homo sapiens hypothetical protein FLJ11273 (FLJ11273), mRNA. /FEA=mRNA /GEN=FLJ11273 /PROD=hypothetical protein FLJ11273 /DB_XREF=gi:8922965 /UG=Hs.3542 hypothetical protein FLJ11273 /FL=gb:NM_018374.1	NM_018374		NP_060844

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218937_at	0.033381	gb:NM_017810.1 /DEF=Homo sapiens hypothetical protein FLJ20417 (FLJ20417), mRNA. /FEA=mRNA /GEN=FLJ20417 /PROD=hypothetical protein FLJ20417 /DB_XREF=gi:8923385 /UG=Hs.10710 hypothetical protein FLJ20417 /FL=gb:NM_017810.1	NM_017810		NP_060280
218941_at	1.26E-04	gb:NM_012164.2 /DEF=Homo sapiens f-box and WD-40 domain protein 2 (FBXW2), mRNA. /FEA=mRNA /GEN=FBXW2 /PROD=f-box and WD-40 domain protein 2 /DB_XREF=gi:7549806 /UG=Hs.13755 f-box and WD-40 domain protein 2 /FL=gb:AF176698.1 gb:NM_012164.2	NM_012164		NP_036296
218942_at	0.002536	gb:NM_024779.1 /DEF=Homo sapiens hypothetical protein FLJ22055 (FLJ22055), mRNA. /FEA=mRNA /GEN=FLJ22055 /PROD=hypothetical protein FLJ22055 /DB_XREF=gi:13376135 /UG=Hs.144502 hypothetical protein FLJ22055 /FL=gb:NM_024779.1	NM_024779		NP_079055
218946_at	0.033381	gb:NM_015700.1 /DEF=Homo sapiens HIRIP5 protein (HIRIP5), mRNA. /FEA=mRNA /GEN=HIRIP5 /PROD=HIRIP5 protein /DB_XREF=gi:7661719 /UG=Hs.17368 HIRIP5 protein; HIRA-interacting protein 5 /FL=gb:AF132967.1 gb:NM_015700.1	NM_015700		NP_056515
218949_s_at	3.26E-04	gb:NM_018292.1 /DEF=Homo sapiens hypothetical protein FLJ10989 (FLJ10989), mRNA. /FEA=mRNA /GEN=FLJ10989 /PROD=hypothetical protein FLJ10989 /DB_XREF=gi:8922807 /UG=Hs.210778 hypothetical protein FLJ10989 /FL=gb:AL136679.1 gb:NM_018292.1	NM_018292		NP_060762

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218961_s_at	0.018693	gb:NM_007254.1 /DEF=Homo sapiens polynucleotide kinase 3-phosphatase (PNKP), mRNA. /FEA=mRNA /GEN=PNKP /PROD=polynucleotide kinase 3-phosphatase /DB_XREF=gi:6005835 /UG=Hs.78016 polynucleotide kinase 3-phosphatase /FL=gb:AF120499.1 gb:AF125807.1 gb:AF126486.1 gb:NM_007254.1	NM_007254		NP_009185
218963_s_at	0.021448	gb:NM_015515.1 /DEF=Homo sapiens DKFZP434G032 protein (DKFZP434G032), mRNA. /FEA=mRNA /GEN=DKFZP434G032 /PROD=DKFZP434G032 protein /DB_XREF=gi:7661573 /UG=Hs.9029 DKFZP434G032 protein /FL=gb:AF102848.1 gb:NM_015515.1	NM_015515		NP_775320
218964_at	0.021312	gb:NM_006465.1 /DEF=Homo sapiens dead ringer (Drosophila)-like 2 (bright and dead ringer) (DRIL2), mRNA. /FEA=mRNA /GEN=DRIL2 /PROD=dead ringer-like 2 /DB_XREF=gi:5453637 /UG=Hs.10431 dead ringer (Drosophila)-like 2 (bright and dead ringer) /FL=gb:AF116846.1 gb:NM_006465.1	NM_006465		NP_006456
218967_s_at	0.003597	gb:NM_030664.1 /DEF=Homo sapiens phosphotriesterase related (PTER), mRNA. /FEA=mRNA /GEN=PTER /PROD=phosphotriesterase related /DB_XREF=gi:13489055 /UG=Hs.129915 phosphotriesterase related /FL=gb:AF212237.1 gb:NM_030664.1	NM_030664		NP_109589

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218968_s_at	0.03018	gb:NM_018197.1 /DEF=Homo sapiens hypothetical protein FLJ10734 (FLJ10734), mRNA. /FEA=mRNA /GEN=FLJ10734 /PROD=hypothetical protein FLJ10734 /DB_XREF=gi:8922626 /UG=Hs.146332 hypothetical protein FLJ10734 /FL=gb:NM_018197.1	NM_018197		NP_071371
218969_at	0.023827	gb:NM_016069.1 /DEF=Homo sapiens CGI-136 protein (LOC51025), mRNA. /FEA=mRNA /GEN=LOC51025 /PROD=CGI-136 protein /DB_XREF=gi:7705633 /UG=Hs.16089 CGI-136 protein /FL=gb:BC005024.1 gb:AF151894.1 gb:NM_016069.1	NM_016069		NP_057153
218970_s_at	0.016934	gb:NM_015960.1 /DEF=Homo sapiens CGI-32 protein (LOC51076), mRNA. /FEA=mRNA /GEN=LOC51076 /PROD=CGI-32 protein /DB_XREF=gi:7705727 /UG=Hs.16606 CGI-32 protein /FL=gb:AF132966.1 gb:NM_015960.1	NM_015960		NP_057044
218979_at	0.001911	gb:NM_024945.1 /DEF=Homo sapiens hypothetical protein FLJ12888 (FLJ12888), mRNA. /FEA=mRNA /GEN=FLJ12888 /PROD=hypothetical protein FLJ12888 /DB_XREF=gi:13376426 /UG=Hs.284137 hypothetical protein FLJ12888 /FL=gb:NM_024945.1	NM_024945		NP_079221
218981_at	1.28E-04	gb:NM_020186.1 /DEF=Homo sapiens DC11 protein (DC11), mRNA. /FEA=mRNA /GEN=DC11 /PROD=DC11 protein /DB_XREF=gi:9910179 /UG=Hs.42785 DC11 protein /FL=gb:AF201933.1 gb:NM_020186.1	NM_020186		NP_064571

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218983_at	1.26E-04	gb:NM_016546.1 /DEF=Homo sapiens complement C1r-like proteinase precursor, (LOC51279), mRNA. /FEA=mRNA /GEN=LOC51279 /PROD=complement C1r-like proteinase precursor, /DB_XREF=gi:7706082 /UG=Hs.98571 complement C1r-like proteinase precursor, /FL=gb:AF178985.1 gb:NM_016546.1	NM_016546		NP_057630
218992_at	1.72E-04	gb:NM_018465.1 /DEF=Homo sapiens uncharacterized hematopoietic stemprogenitor cells protein MDS030 (MDS030), mRNA. /FEA=mRNA /GEN=MDS030 /PROD=uncharacterized hematopoietic stemprogenitorcells protein MDS030 /DB_XREF=gi:8923931 /UG=Hs.181385 uncharacterized hematopoietic stemprogenitor cells protein MDS030 /FL=gb:AF220050.1 gb:NM_018465.1 gb:AF225420.1	NM_018465		NP_060935
218996_at	0.001786	gb:NM_013342.1 /DEF=Homo sapiens TCF3 (E2A) fusion partner (in childhood Leukemia) (TFPT), mRNA. /FEA=mRNA /GEN=TFPT /PROD=TCF3 (E2A) fusion partner (in childhoodLeukemia) /DB_XREF=gi:7019370 /UG=Hs.233765 TCF3 (E2A) fusion partner (in childhood Leukemia) /FL=gb:BC001728.1 gb:BC004281.1 gb:AF052052.1 gb:NM_013342.1	NM_013342		NP_037474

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218997_at	0.006705	gb:NM_022490.1 /DEF=Homo sapiens hypothetical protein FLJ13390 similar to PAF53 (FLJ13390), mRNA. /FEA=mRNA /GEN=FLJ13390 /PROD=hypothetical protein FLJ13390 similar to PAF53 /DB_XREF=gi:11968046 /UG=Hs.24884 hypothetical protein FLJ13390 similar to PAF53 /FL=gb:NM_022490.1 gb:BC001337.1	NM_022490		NP_071935
218998_at	1.26E-04	gb:NM_017832.1 /DEF=Homo sapiens hypothetical protein FLJ20457 (FLJ20457), mRNA. /FEA=mRNA /GEN=FLJ20457 /PROD=hypothetical protein FLJ20457 /DB_XREF=gi:8923430 /UG=Hs.29276 hypothetical protein FLJ20457 /FL=gb:AF153417.1 gb:NM_017832.1	NM_017832		NP_060302
219001_s_at	0.041055	gb:NM_024345.1 /DEF=Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA. /FEA=mRNA /GEN=MGC10765 /PROD=hypothetical protein MGC10765 /DB_XREF=gi:13236596 /UG=Hs.317589 hypothetical protein MGC10765 /FL=gb:BC003520.1 gb:NM_024345.1	NM_024345		NP_077321
219004_s_at	0.007581	gb:NM_018944.1 /DEF=Homo sapiens chromosome 21 open reading frame 45 (C21ORF45), mRNA. /FEA=mRNA /GEN=C21ORF45 /PROD=chromosome 21 open reading frame 45 /DB_XREF=gi:9506436 /UG=Hs.49932 chromosome 21 open reading frame 45 /FL=gb:NM_018944.1	NM_018944		NP_061817

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219012_s_at	0.006966	gb:NM_020193.1 /DEF=Homo sapiens GL002 protein (GL002), mRNA. /FEA=mRNA /GEN=GL002 /PROD=GL002 protein /DB_XREF=gi:9910245 /UG=Hs.9567 GL002 protein /FL=gb:AF226047.1 gb:NM_020193.1	NM_020193		NP_064578
219014_at	1.26E-04	gb:NM_016619.1 /DEF=Homo sapiens hypothetical protein (LOC51316), mRNA. /FEA=mRNA /GEN=LOC51316 /PROD=hypothetical protein /DB_XREF=gi:7706157 /UG=Hs.107139 hypothetical protein /FL=gb:AF208846.1 gb:NM_016619.1	NM_016619		NP_057703
219015_s_at	0.00228	gb:NM_018466.1 /DEF=Homo sapiens uncharacterized hematopoietic stemprogenitor cells protein MDS031 (MDS031), mRNA. /FEA=mRNA /GEN=MDS031 /PROD=uncharacterized hematopoietic stemprogenitorcells protein MDS031 /DB_XREF=gi:8923933 /UG=Hs.110853 uncharacterized hematopoietic stemprogenitor cells protein MDS031 /FL=gb:BC005336.1 gb:AF220051.1 gb:NM_018466.1	NM_018466		NP_060936
219016_at	1.36E-04	gb:NM_021826.1 /DEF=Homo sapiens hypothetical protein FLJ13149 (FLJ13149), mRNA. /FEA=mRNA /GEN=FLJ13149 /PROD=hypothetical protein FLJ13149 /DB_XREF=gi:11141902 /UG=Hs.112188 hypothetical protein FLJ13149 /FL=gb:NM_021826.1	NM_021826		NP_068598

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219017_at	1.26E-04	gb:NM_018638.2 /DEF=Homo sapiens ethanolamine kinase (EKI1), mRNA. /FEA=mRNA /GEN=EKI1 /PROD=ethanolamine kinase /DB_XREF=gi:10092614 /UG=Hs.120439 ethanolamine kinase /FL=gb:NM_018638.2 gb:AF207600.2	NM_018638		NP_061108
219020_at	0.027418	gb:NM_022460.1 /DEF=Homo sapiens hypothetical protein FLJ14249 similar to HS1 binding protein 3 (FLJ14249), mRNA. /FEA=mRNA /GEN=FLJ14249 /PROD=hypothetical protein FLJ14249 similar to HS1 binding protein 3 /DB_XREF=gi:11968000 /UG=Hs.125520 hypothetical protein FLJ14249 similar to HS1 binding protein 3 /FL=gb:NM_022460.1	NM_022460		NP_567825
219035_s_at	1.26E-04	gb:NM_025126.1 /DEF=Homo sapiens hypothetical protein FLJ21786 (FLJ21786), mRNA. /FEA=mRNA /GEN=FLJ21786 /PROD=hypothetical protein FLJ21786 /DB_XREF=gi:13376704 /UG=Hs.316809 hypothetical protein FLJ21786 /FL=gb:NM_025126.1	NM_025126		NP_079402
219049_at	1.26E-04	gb:NM_018371.1 /DEF=Homo sapiens hypothetical protein FLJ11264 (FLJ11264), mRNA. /FEA=mRNA /GEN=FLJ11264 /PROD=hypothetical protein FLJ11264 /DB_XREF=gi:8922959 /UG=Hs.11260 hypothetical protein FLJ11264 /FL=gb:NM_018371.1	NM_018371		NP_060841

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219065_s_at	0.010205	gb:NM_015955.1 /DEF=Homo sapiens CGI-27 protein (LOC51072), mRNA. /FEA=mRNA /GEN=LOC51072 /PROD=CGI-27 protein /DB_XREF=gi:7705719 /UG=Hs.20814 CGI-27 protein /FL=gb:AF132961.1 gb:NM_015955.1	NM_015955		NP_057039
219070_s_at	0.001483	gb:NM_023948.1 /DEF=Homo sapiens hypothetical protein AF053356_CDS3 (AF053356_CDS3), mRNA. /FEA=mRNA /GEN=AF053356_CDS3 /PROD=AF053356_CDS3 protein /DB_XREF=gi:13122607 /UG=Hs.296200 hypothetical protein AF053356_CDS3 /FL=gb:NM_023948.1 gb:BC005042.1	NM_023948		NP_076438
219074_at	0.036688	gb:NM_018241.1 /DEF=Homo sapiens hypothetical protein FLJ10846 (FLJ10846), mRNA. /FEA=mRNA /GEN=FLJ10846 /PROD=hypothetical protein FLJ10846 /DB_XREF=gi:8922706 /UG=Hs.32271 hypothetical protein FLJ10846 /FL=gb:NM_018241.1	NM_018241		NP_060711
219079_at	1.26E-04	gb:NM_016230.1 /DEF=Homo sapiens flavohemoprotein b5+b5R (LOC51167), mRNA. /FEA=mRNA /GEN=LOC51167 /PROD=flavohemoprotein b5+b5R /DB_XREF=gi:7705898 /UG=Hs.5741 flavohemoprotein b5+b5R /FL=gb:AF169803.1 gb:NM_016230.1	NM_016230		NP_057314
219099_at	0.010205	gb:NM_020375.1 /DEF=Homo sapiens chromosome 12 open reading frame 5 (C12ORF5), mRNA. /FEA=mRNA /GEN=C12ORF5 /PROD=chromosome 12 open reading frame 5 /DB_XREF=gi:9966848 /UG=Hs.24792 chromosome 12 open reading frame 5 /FL=gb:NM_020375.1	NM_020375		NP_065108

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219104_at	5.03E-04	gb:NM_016422.1 /DEF=Homo sapiens C3HC4-like zinc finger protein (ZFP26), mRNA. /FEA=mRNA /GEN=ZFP26 /PROD=C3HC4-like zinc finger protein /DB_XREF=gi:7706776 /UG=Hs.44685 C3HC4-like zinc finger protein. /FL=gb:AF214680.1 gb:NM_016422.1	NM_016422		NP_057506
219112_at	0.00161	gb:NM_016340.1 /DEF=Homo sapiens PDZ domain-containing guanine nucleotide exchange factor I (LOC51735), mRNA. /FEA=mRNA /GEN=LOC51735 /PROD=PDZ domain-containing guanine nucleotideexchange factor I /DB_XREF=gi:7706512 /UG=Hs.174795 PDZ domain-containing guanine nucleotide exchange factor I /FL=gb:AF117947.1 gb:NM_016340.1	NM_016340		NP_057424
219116_s_at	0.003395	gb:NM_018185.1 /DEF=Homo sapiens hypothetical protein FLJ10704 (FLJ10704), mRNA. /FEA=mRNA /GEN=FLJ10704 /PROD=hypothetical protein FLJ10704 /DB_XREF=gi:8922602 /UG=Hs.23618 hypothetical protein FLJ10704 /FL=gb:NM_018185.1	NM_018185		NP_060655
219123_at	0.022926	gb:NM_014519.1 /DEF=Homo sapiens zinc finger protein 232 (ZNF232), mRNA. /FEA=mRNA /GEN=ZNF232 /PROD=zinc finger protein 232 /DB_XREF=gi:7657704 /UG=Hs.279914 zinc finger protein 232 /FL=gb:NM_014519.1	NM_014519		NP_055334

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219129_s_at	1.49E-04	gb:NM_024632.1 /DEF=Homo sapiens hypothetical protein FLJ11526 (FLJ11526), mRNA. /FEA=mRNA /GEN=FLJ11526 /PROD=hypothetical protein FLJ11526 /DB_XREF=gi:13375861 /UG=Hs.38750 hypothetical protein FLJ11526 /FL=gb:NM_024632.1	NM_024632		NP_078908
219132_at	5.03E-04	gb:NM_021255.1 /DEF=Homo sapiens pellino (Drosophila) homolog 2 (PELI2), mRNA. /FEA=mRNA /GEN=PELI2 /PROD=pellino (Drosophila) homolog 2 /DB_XREF=gi:10864062 /UG=Hs.44038 pellino (Drosophila) homolog 2 /FL=gb:AF302502.1 gb:NM_021255.1	NM_021255		NP_067078
219137_s_at	1.26E-04	gb:NM_020194.1 /DEF=Homo sapiens GL004 protein (GL004), mRNA. /FEA=mRNA /GEN=GL004 /PROD=GL004 protein /DB_XREF=gi:9910247 /UG=Hs.7045 GL004 protein /FL=gb:AF226049.1 gb:NM_020194.1	NM_020194		NP_064579
219149_x_at	0.004908	gb:NM_016216.1 /DEF=Homo sapiens RNA lariat debranching enzyme (LOC51163), mRNA. /FEA=mRNA /GEN=LOC51163 /PROD=RNA lariat debranching enzyme /DB_XREF=gi:7705890 /UG=Hs.118194 RNA lariat debranching enzyme /FL=gb:AF180919.2 gb:NM_016216.1	NM_016216		NP_057300
219156_at	1.26E-04	gb:NM_018373.1 /DEF=Homo sapiens hypothetical protein FLJ11271 (FLJ11271), mRNA. /FEA=mRNA /GEN=FLJ11271 /PROD=hypothetical protein FLJ11271 /DB_XREF=gi:8922963 /UG=Hs.109654 hypothetical protein FLJ11271 /FL=gb:NM_018373.1	NM_018373		NP_060843

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219157_at	0.001116	gb:NM_007246.1 /DEF=Homo sapiens kelch (Drosophila)-like 2 (Mayven) (KLHL2), mRNA. /FEA=mRNA /GEN=KLHL2 /PROD=kelch (Drosophila)-like 2 (Mayven) /DB_XREF=gi:6005702 /UG=Hs.122967 kelch (Drosophila)-like 2 (Mayven) /FL=gb:AF059569.1 gb:NM_007246.1	NM_007246		NP_009177
219158_s_at	6.68E-04	gb:NM_025085.1 /DEF=Homo sapiens hypothetical protein FLJ13340 (FLJ13340), mRNA. /FEA=mRNA /GEN=FLJ13340 /PROD=hypothetical protein FLJ13340 /DB_XREF=gi:13376639 /UG=Hs.125034 hypothetical protein FLJ13340 /FL=gb:AF327722.1 gb:NM_025085.1	NM_025085		NP_476516
219161_s_at	0.004341	gb:NM_016951.2 /DEF=Homo sapiens transmembrane proteolipid (HSPC224), mRNA. /FEA=mRNA /GEN=HSPC224 /PROD=transmembrane proteolipid /DB_XREF=gi:10092593 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM_016951.2 gb:BC004380.1 gb:AF057306.1 gb:AF151058.1 gb:AF135380.2	NM_016951		NP_058647
219163_at	0.007052	gb:NM_017656.1 /DEF=Homo sapiens hypothetical protein FLJ20079 (FLJ20079), mRNA. /FEA=mRNA /GEN=FLJ20079 /PROD=hypothetical protein FLJ20079 /DB_XREF=gi:8923085 /UG=Hs.165948 hypothetical protein FLJ20079 /FL=gb:NM_017656.1	NM_017656		NP_060126

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219165_at	0.016934	gb:NM_021630.1 /DEF=Homo sapiens PDZ-LIM protein mystique (LOC59346), mRNA. /FEA=mRNA /GEN=LOC59346 /PROD=PDZ-LIM protein mystique /DB_XREF=gi:11055999 /UG=Hs.19447 PDZ-LIM protein mystique /FL=gb:AY007729.1 gb:NM_021630.1	NM_021630		NP_789847
219172_at	0.034221	gb:NM_024954.1 /DEF=Homo sapiens hypothetical protein FLJ11807 (FLJ11807), mRNA. /FEA=mRNA /GEN=FLJ11807 /PROD=hypothetical protein FLJ11807 /DB_XREF=gi:13376438 /UG=Hs.285813 hypothetical protein FLJ11807 /FL=gb:NM_024954.1	NM_024954		NP_079230
219180_s_at	0.02145	hypothetical protein FLJ20695	AI817074	Hs.5420	NP_060399
219190_s_at	0.005852	gb:NM_017629.1 /DEF=Homo sapiens hypothetical protein FLJ20033 (FLJ20033), mRNA. /FEA=mRNA /GEN=FLJ20033 /PROD=hypothetical protein FLJ20033 /DB_XREF=gi:8923033 /UG=Hs.134757 hypothetical protein FLJ20033 /FL=gb:NM_017629.1	NM_017629		NP_060099
219191_s_at	3.26E-04	gb:NM_016293.1 /DEF=Homo sapiens bridging integrator 2 (BIN2), mRNA. /FEA=mRNA /GEN=BIN2 /PROD=breast cancer associated protein BRAP1 /DB_XREF=gi:7706486 /UG=Hs.14770 bridging integrator 2 /FL=gb:AF146531.1 gb:AB032698.1 gb:NM_016187.1 gb:NM_016293.1	NM_016293		NP_057377
219221_at	0.003539	gb:NM_024724.1 /DEF=Homo sapiens hypothetical protein FLJ22332 (FLJ22332), mRNA. /FEA=mRNA /GEN=FLJ22332 /PROD=hypothetical protein FLJ22332 /DB_XREF=gi:13376033 /UG=Hs.111092 hypothetical protein FLJ22332 /FL=gb:NM_024724.1	NM_024724		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219228_at	6.68E-04	gb:NM_018555.2 /DEF=Homo sapiens C2H2-like zinc finger protein (ZNF361), mRNA. /FEA=mRNA /GEN=ZNF361 /PROD=C2H2-like zinc finger protein /DB_XREF=gi:10092612 /UG=Hs.147644 zinc finger protein 331 /FL=gb:AF251515.2 gb:NM_018555.2 gb:AF272148.1	NM_018555		NP_061025
219229_at	0.001116	gb:NM_013272.2 /DEF=Homo sapiens solute carrier family 21 (organic anion transporter), member 11 (SLC21A11), mRNA. /FEA=mRNA /GEN=SLC21A11 /PROD=solute carrier family 21 (organic anion transporter), member 11 /DB_XREF=gi:7706713 /UG=Hs.14805 solute carrier family 21 (organic anion transporter), member 11 /FL=gb:AF205074.1 gb:AF187816.1 gb:AB031050.2 gb:NM_013272.2	NM_013272		NP_037404
219232_s_at	0.014208	gb:NM_022073.1 /DEF=Homo sapiens hypothetical protein FLJ21620 (FLJ21620), mRNA. /FEA=mRNA /GEN=FLJ21620 /PROD=hypothetical protein FLJ21620 /DB_XREF=gi:11545786 /UG=Hs.18878 hypothetical protein FLJ21620 /FL=gb:NM_022073.1	NM_022073		NP_203130
219238_at	0.006966	gb:NM_017837.1 /DEF=Homo sapiens hypothetical protein FLJ20477 (FLJ20477), mRNA. /FEA=mRNA /GEN=FLJ20477 /PROD=hypothetical protein FLJ20477 /DB_XREF=gi:8923441 /UG=Hs.26994 hypothetical protein FLJ20477 /FL=gb:NM_017837.1	NM_017837		NP_060307

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219253_at	1.26E-04	gb:NM_024121.1 /DEF=Homo sapiens hypothetical protein FLJ20979 (FLJ20979), mRNA. /FEA=mRNA /GEN=FLJ20979 /PROD=hypothetical protein FLJ20979 /DB_XREF=gi:13129145 /UG=Hs.44680 hypothetical protein FLJ20979 /FL=gb:NM_024121.1	NM_024121		
219259_at	0.00228	gb:NM_022367.1 /DEF=Homo sapiens hypothetical protein FLJ12287 similar to semaphorins (FLJ12287), mRNA. /FEA=mRNA /GEN=FLJ12287 /PROD=hypothetical protein FLJ12287 similar to semaphorins /DB_XREF=gi:11641290 /UG=Hs.7634 hypothetical protein FLJ12287 similar to semaphorins /FL=gb:NM_022367.1 gb:AB029394.1	NM_022367		NP_071762
219266_at	0.02145	gb:NM_021632.1 /DEF=Homo sapiens zinc-finger protein ZBRK1 (ZBRK1), mRNA. /FEA=mRNA /GEN=ZBRK1 /PROD=zinc-finger protein ZBRK1 /DB_XREF=gi:11056003 /UG=Hs.130965 zinc-finger protein ZBRK1 /FL=gb:AF295096.1 gb:AF309561.1 gb:NM_021632.1	NM_021632		NP_067645
219281_at	1.26E-04	gb:NM_012331.2 /DEF=Homo sapiens methionine sulfoxide reductase A (MSRA), mRNA. /FEA=mRNA /GEN=MSRA /PROD=methionine sulfoxide reductase A /DB_XREF=gi:13259538 /UG=Hs.26458 methionine sulfoxide reductase A /FL=gb:NM_012331.2 gb:AF183420.1	NM_012331		NP_036463

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219283_at	1.26E-04	gb:NM_014158.1 /DEF=Homo sapiens HSPC067 protein (HSPC067), mRNA. /FEA=mRNA /GEN=HSPC067 /PROD=HSPC067 protein /DB_XREF=gi:7661771 /UG=Hs.279938 HSPC067 protein /FL=gb:AF161552.1 gb:NM_014158.1	NM_014158		
219292_at	3.56E-04	gb:NM_018105.1 /DEF=Homo sapiens hypothetical protein FLJ10477 (FLJ10477), mRNA. /FEA=mRNA /GEN=FLJ10477 /PROD=hypothetical protein FLJ10477 /DB_XREF=gi:8922445 /UG=Hs.7432 hypothetical protein FLJ10477 /FL=gb:NM_018105.1	NM_018105		NP_060575
219297_at	0.026891	gb:NM_019045.1 /DEF=Homo sapiens similar to rab11-binding protein (FLJ11116), mRNA. /FEA=mRNA /GEN=FLJ11116 /PROD=similar to rab11-binding protein /DB_XREF=gi:9506636 /UG=Hs.98510 similar to rab11-binding protein /FL=gb:NM_019045.1	NM_019045		NP_061918
219304_s_at	0.041055	gb:NM_025208.1 /DEF=Homo sapiens spinal cord-derived growth factor-B (SCDGF-B), mRNA. /FEA=mRNA /GEN=SCDGF-B /PROD=spinal cord-derived growth factor-B /DB_XREF=gi:13376807 /UG=Hs.112885 spinal cord-derived growth factor-B /FL=gb:AB033832.1 gb:AF113216.1 gb:NM_025208.1 gb:AY027517.1	NM_025208		NP_149126
219322_s_at	0.02145	gb:NM_017818.1 /DEF=Homo sapiens WD repeat domain 8 (WDR8), mRNA. /FEA=mRNA /GEN=WDR8 /PROD=WD repeat domain 8 /DB_XREF=gi:8923402 /UG=Hs.31714 hypothetical protein FLJ20430 /FL=gb:AB034912.1 gb:NM_017818.1	NM_017818		NP_060288

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219324_at	0.001173	gb:NM_024313.1 /DEF=Homo sapiens hypothetical protein MGC3731 (MGC3731), mRNA. /FEA=mRNA /GEN=MGC3731 /PROD=hypothetical protein MGC3731 /DB_XREF=gi:13236552 /UG=Hs.326372 hypothetical protein MGC3731 /FL=gb:BC002808.1 gb:NM_024313.1	NM_024313		NP_077289
219326_s_at	3.93E-04	gb:NM_006577.2 /DEF=Homo sapiens beta-1,3-N-acetylglucosaminyltransferase (BETA3GNT), mRNA. /FEA=mRNA /GEN=BETA3GNT /PROD=beta-1,3-N-acetylglucosaminyltransferase /DB_XREF=gi:9845237 /UG=Hs.48730 beta-1,3-N-acetylglucosaminyltransferase bGnT-2 /FL=gb:AF288208.1 gb:AF092051.2 gb:NM_006577.2	NM_006577		NP_150274
219329_s_at	0.007779	gb:NM_016085.1 /DEF=Homo sapiens apoptosis related protein APR-3 (APR-3), mRNA. /FEA=mRNA /GEN=APR-3 /PROD=apoptosis related protein APR-3 /DB_XREF=gi:7706360 /UG=Hs.9527 apoptosis related protein APR-3 /FL=gb:AF144055.2 gb:NM_016085.1	NM_016085		NP_542159
219336_s_at	0.034317	gb:NM_015947.1 /DEF=Homo sapiens CGI-18 protein (LOC51008), mRNA. /FEA=mRNA /GEN=LOC51008 /PROD=CGI-18 protein /DB_XREF=gi:7705601 /UG=Hs.121599 CGI-18 protein /FL=gb:AY013290.1 gb:AF132952.1 gb:NM_015947.1	NM_015947		NP_057031

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219356_s_at	0.001116	gb:NM_016410.1 /DEF=Homo sapiens hypothetical protein (HSPC177), mRNA. /FEA=mRNA /GEN=HSPC177 /PROD=hypothetical protein /DB_XREF=gi:7705488 /UG=Hs.279777 hypothetical protein /FL=gb:AF161525.1 gb:NM_016410.1	NM_016410		NP_057494
219357_at	0.007558	gb:NM_014027.1 /DEF=Homo sapiens HSPC018 protein (HSPC018), mRNA. /FEA=mRNA /GEN=HSPC018 /PROD=HSPC018 protein /DB_XREF=gi:7661735 /UG=Hs.279853 HSPC018 protein /FL=gb:BC001701.1 gb:AF077204.1 gb:NM_014027.1	NM_014027		NP_054746
219359_at	3.26E-04	gb:NM_025092.1 /DEF=Homo sapiens hypothetical protein FLJ22635 (FLJ22635), mRNA. /FEA=mRNA /GEN=FLJ22635 /PROD=hypothetical protein FLJ22635 /DB_XREF=gi:13376651 /UG=Hs.288529 hypothetical protein FLJ22635 /FL=gb:NM_025092.1	NM_025092		NP_079368
219371_s_at	0.005852	gb:NM_016270.1 /DEF=Homo sapiens Kruppel-like factor (LOC51713), mRNA. /FEA=mRNA /GEN=LOC51713 /PROD=Kruppel-like factor /DB_XREF=gi:7706468 /UG=Hs.107740 Kruppel-like factor 2 (lung) /FL=gb:AF134053.1 gb:AF205849.1 gb:NM_016270.1	NM_016270		NP_057354
219384_s_at	1.72E-04	gb:NM_012091.2 /DEF=Homo sapiens adenosine deaminase, tRNA-specific 1 (ADAT1), mRNA. /FEA=mRNA /GEN=ADAT1 /PROD=adenosine deaminase, tRNA-specific 1 /DB_XREF=gi:7382475 /UG=Hs.188661 adenosine deaminase, tRNA-specific 1 /FL=gb:BC002758.1 gb:AF125188.1 gb:NM_012091.2	NM_012091		NP_036223

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219394_at	0.00161	gb:NM_024419.1 /DEF=Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA. /FEA=mRNA /GEN=PGS1 /PROD=Phosphatidylglycerophosphate Synthase /DB_XREF=gi:13259369 /UG=Hs.278682 Phosphatidylglycerophosphate Synthase /FL=gb:NM_024419.1	NM_024419		NP_077733
219396_s_at	0.021312	gb:NM_024608.1 /DEF=Homo sapiens hypothetical protein FLJ22402 (FLJ22402), mRNA. /FEA=mRNA /GEN=FLJ22402 /PROD=hypothetical protein FLJ22402 /DB_XREF=gi:13375816 /UG=Hs.28355 hypothetical protein FLJ22402 /FL=gb:NM_024608.1	NM_024608		NP_078884
219397_at	0.003171	gb:NM_025147.1 /DEF=Homo sapiens hypothetical protein FLJ13448 (FLJ13448), mRNA. /FEA=mRNA /GEN=FLJ13448 /PROD=hypothetical protein FLJ13448 /DB_XREF=gi:13376736 /UG=Hs.288945 hypothetical protein FLJ13448 /FL=gb:NM_025147.1	NM_025147		NP_079423
219403_s_at	1.41E-04	gb:NM_006665.1 /DEF=Homo sapiens heparanase (HPSE), mRNA. /FEA=mRNA /GEN=HPSE /PROD=heparanase /DB_XREF=gi:5729872 /UG=Hs.44227 heparanase /FL=gb:AF165154.1 gb:AF152376.1 gb:NM_006665.1 gb:AF084467.1 gb:AF155510.1	NM_006665		NP_006656
219405_at	0.023856	gb:NM_018073.1 /DEF=Homo sapiens hypothetical protein FLJ10369 (FLJ10369), mRNA. /FEA=mRNA /GEN=FLJ10369 /PROD=hypothetical protein FLJ10369 /DB_XREF=gi:8922379 /UG=Hs.53940 hypothetical protein FLJ10369 /FL=gb:NM_018073.1	NM_018073		NP_060543

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219409_at	0.046189	gb:NM_024700.1 /DEF=Homo sapiens hypothetical protein FLJ12553 (FLJ12553), mRNA. /FEA=mRNA /GEN=FLJ12553 /PROD=hypothetical protein FLJ12553 /DB_XREF=gi:13375986 /UG=Hs.99601 hypothetical protein FLJ12553 /FL=gb:NM_024700.1	NM_024700		NP_078976
219410_at	0.037692	gb:NM_018004.1 /DEF=Homo sapiens hypothetical protein FLJ10134 (FLJ10134), mRNA. /FEA=mRNA /GEN=FLJ10134 /PROD=hypothetical protein FLJ10134 /DB_XREF=gi:8922242 /UG=Hs.104800 hypothetical protein FLJ10134 /FL=gb:NM_018004.1	NM_018004		NP_060474
219426_at	7.58E-04	gb:NM_024852.1 /DEF=Homo sapiens hypothetical protein FLJ12765 (FLJ12765), mRNA. /FEA=mRNA /GEN=FLJ12765 /PROD=hypothetical protein FLJ12765 /DB_XREF=gi:13376274 /UG=Hs.189829 hypothetical protein FLJ12765 /FL=gb:NM_024852.1	NM_024852		NP_803171
219444_at	1.26E-04	gb:NM_021946.1 /DEF=Homo sapiens hypothetical protein FLJ11362 (FLJ11362), mRNA. /FEA=mRNA /GEN=FLJ11362 /PROD=hypothetical protein FLJ11362 /DB_XREF=gi:11345489 /UG=Hs.8929 hypothetical protein FLJ11362 /FL=gb:NM_021946.1	NM_021946		NP_068765

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219445_at	1.49E-04	gb:NM_015711.1 /DEF=Homo sapiens glioma tumor suppressor candidate region gene 1 (GLTSCR1), mRNA. /FEA=mRNA /GEN=GLTSCR1 /PROD=glioma tumor suppressor candidate region gene 1 /DB_XREF=gi:7657127 /UG=Hs.97244 glioma tumor suppressor candidate region gene 1 /FL=gb:AF182077.1 gb:NM_015711.1	NM_015711		NP_056526
219446_at	0.042466	gb:NM_018157.1 /DEF=Homo sapiens hypothetical protein FLJ10620 (FLJ10620), mRNA. /FEA=mRNA /GEN=FLJ10620 /PROD=hypothetical protein FLJ10620 /DB_XREF=gi:8922554 /UG=Hs.99445 hypothetical protein FLJ10620 /FL=gb:NM_018157.1	NM_018157		NP_060627
219449_s_at	3.26E-04	gb:NM_017866.1 /DEF=Homo sapiens hypothetical protein FLJ20533 (FLJ20533), mRNA. /FEA=mRNA /GEN=FLJ20533 /PROD=hypothetical protein FLJ20533 /DB_XREF=gi:8923499 /UG=Hs.106650 hypothetical protein FLJ20533 /FL=gb:BC002748.1 gb:NM_017866.1	NM_017866		NP_060336
219457_s_at	0.005039	gb:NM_024832.1 /DEF=Homo sapiens hypothetical protein FLJ22439 (FLJ22439), mRNA. /FEA=mRNA /GEN=FLJ22439 /PROD=hypothetical protein FLJ22439 /DB_XREF=gi:13376237 /UG=Hs.180040 hypothetical protein FLJ22439 /FL=gb:NM_024832.1	NM_024832		NP_079108

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219458_s_at	0.026891	gb:NM_022072.1 /DEF=Homo sapiens hypothetical protein FLJ22609 (FLJ22609), mRNA. /FEA=mRNA /GEN=FLJ22609 /PROD=hypothetical protein FLJ22609 /DB_XREF=gi:11545784 /UG=Hs.18740 hypothetical protein FLJ22609 /FL=gb:NM_022072.1	NM_022072		NP_071355
219459_at	0.03002	gb:NM_018082.1 /DEF=Homo sapiens hypothetical protein FLJ10388 (FLJ10388), mRNA. /FEA=mRNA /GEN=FLJ10388 /PROD=hypothetical protein FLJ10388 /DB_XREF=gi:8922398 /UG=Hs.197642 hypothetical protein FLJ10388 /FL=gb:NM_018082.1	NM_018082		NP_060552
219460_s_at	1.28E-04	gb:NM_017849.1 /DEF=Homo sapiens hypothetical protein FLJ20507 (FLJ20507), mRNA. /FEA=mRNA /GEN=FLJ20507 /PROD=hypothetical protein FLJ20507 /DB_XREF=gi:8923465 /UG=Hs.202955 hypothetical protein FLJ20507 /FL=gb:NM_017849.1	NM_017849		NP_060319
219467_at	0.033381	gb:NM_017676.1 /DEF=Homo sapiens hypothetical protein FLJ20125 (FLJ20125), mRNA. /FEA=mRNA /GEN=FLJ20125 /PROD=hypothetical protein FLJ20125 /DB_XREF=gi:8923123 /UG=Hs.24088 hypothetical protein FLJ20125 /FL=gb:NM_017676.1	NM_017676		NP_060146

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219485_s_at	0.001116	gb:NM_002814.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 10 (PSMD10), mRNA. /FEA=mRNA /GEN=PSMD10 /PROD=proteasome (prosome, macropain) 26S subunit, non-ATPase, 10 /DB_XREF=gi:4506216 /UG=Hs.7756 proteasome (prosome, macropain) 26S subunit, non-ATPase, 10 /FL=gb:AB009619.1 gb:D83197.1 gb:NM_002814.1	NM_002814		NP_736606
219492_at	3.26E-04	gb:NM_012110.1 /DEF=Homo sapiens cystein-rich hydrophobic domain 2 (CHIC2), mRNA. /FEA=mRNA /GEN=CHIC2 /PROD=cystein-rich hydrophobic domain 2 /DB_XREF=gi:6912275 /UG=Hs.119488 cystein-rich hydrophobic domain 2 /FL=gb:AF159423.1 gb:NM_012110.1	NM_012110		NP_036242
219493_at	0.005379	gb:NM_024745.1 /DEF=Homo sapiens hypothetical protein FLJ22009 (FLJ22009), mRNA. /FEA=mRNA /GEN=FLJ22009 /PROD=hypothetical protein FLJ22009 /DB_XREF=gi:13376069 /UG=Hs.123253 hypothetical protein FLJ22009 /FL=gb:NM_024745.1	NM_024745		NP_079021
219495_s_at	0.046329	gb:NM_013256.1 /DEF=Homo sapiens zinc finger protein 180 (HHZ168) (ZNF180), mRNA. /FEA=mRNA /GEN=ZNF180 /PROD=zinc finger protein 180 (HHZ168) /DB_XREF=gi:7019578 /UG=Hs.130683 zinc finger protein 180 (HHZ168) /FL=gb:AF192913.1 gb:NM_013256.1	NM_013256		NP_037388

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219497_s_at	0.013088	gb:NM_022893.1 /DEF=Homo sapiens B-cell CLLlymphoma 11A (zinc finger protein) (BCL11A), mRNA. /FEA=mRNA /GEN=BCL11A /PROD=B-cell CLLlymphoma 11A (zinc finger protein) /DB_XREF=gi:13375629 /UG=Hs.130881 B-cell CLLlymphoma 11A (zinc finger protein) /FL=gb:NM_022893.1 gb:NM_018014.1	NM_022893		NP_612569
219502_at	0.046189	gb:NM_018248.1 /DEF=Homo sapiens hypothetical protein FLJ10858 (FLJ10858), mRNA. /FEA=mRNA /GEN=FLJ10858 /PROD=hypothetical protein FLJ10858 /DB_XREF=gi:8922721 /UG=Hs.134403 hypothetical protein FLJ10858 /FL=gb:NM_018248.1	NM_018248		NP_060718
219515_at	0.013223	gb:NM_020228.1 /DEF=Homo sapiens PR domain containing 10 (PRDM10), mRNA. /FEA=mRNA /GEN=PRDM10 /PROD=PR domain containing 10 /DB_XREF=gi:9910503 /UG=Hs.275086 PR domain containing 10 /FL=gb:AF275817.1 gb:NM_020228.1	NM_020228		NP_064613
219520_s_at	0.033381	gb:NM_018458.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM042 (BM042), mRNA. /FEA=mRNA /GEN=BM042 /PROD=uncharacterized bone marrow protein BM042 /DB_XREF=gi:8922101 /UG=Hs.324136 uncharacterized bone marrow protein BM042 /FL=gb:AF217518.1 gb:NM_018458.1	NM_018458		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219528_s_at	0.005852	gb:NM_022898.1 /DEF=Homo sapiens B-cell lymphomaleukaemia 11B (BCL11B), mRNA. /FEA=mRNA /GEN=BCL11B /PROD=B-cell lymphomaleukaemia 11B /DB_XREF=gi:12597634 /UG=Hs.57987 B-cell CLLlymphoma 11B (zinc finger protein) /FL=gb:NM_022898.1	NM_022898		NP_612808
219544_at	0.00228	gb:NM_024808.1 /DEF=Homo sapiens hypothetical protein FLJ22624 (FLJ22624), mRNA. /FEA=mRNA /GEN=FLJ22624 /PROD=hypothetical protein FLJ22624 /DB_XREF=gi:13376190 /UG=Hs.166425 hypothetical protein FLJ22624 /FL=gb:NM_024808.1	NM_024808		NP_079084
219547_at	1.26E-04	gb:NM_004376.1 /DEF=Homo sapiens COX15 (yeast) homolog, cytochrome c oxidase assembly protein (COX15), mRNA. /FEA=mRNA /GEN=COX15 /PROD=COX15 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:4758035 /UG=Hs.226581 COX15 (yeast) homolog, cytochrome c oxidase assembly protein /FL=gb:AF044323.1 gb:NM_004376.1	NM_004376		NP_510870
219549_s_at	7.58E-04	gb:NM_006054.1 /DEF=Homo sapiens reticulon 3 (RTN3), mRNA. /FEA=mRNA /GEN=RTN3 /PROD=reticulon 3 /DB_XREF=gi:5174654 /UG=Hs.252831 reticulon 3 /FL=gb:BC000634.1 gb:AF059524.1 gb:AF119297.1 gb:NM_006054.1	NM_006054		NP_006045

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219551_at	1.32E-04	gb:NM_018456.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM040 (BM040), mRNA. /FEA=mRNA /GEN=BM040 /PROD=uncharacterized bone marrow protein BM040 /DB_XREF=gi:8922098 /UG=Hs.26892 uncharacterized bone marrow protein BM040 /FL=gb:AF217516.1 gb:NM_018456.1	NM_018456		NP_060926
219563_at	0.004341	gb:NM_024633.1 /DEF=Homo sapiens hypothetical protein FLJ21276 (FLJ21276), mRNA. /FEA=mRNA /GEN=FLJ21276 /PROD=hypothetical protein FLJ21276 /DB_XREF=gi:13375863 /UG=Hs.41502 hypothetical protein FLJ21276 /FL=gb:NM_024633.1	NM_024633		NP_078909
219571_s_at	2.05E-04	gb:NM_016265.1 /DEF=Homo sapiens GIOT-3 for gonadotropin inducible transcription repressor-3 (GIOT-3), mRNA. /FEA=mRNA /GEN=GIOT-3 /PROD=GIOT-3 for gonadotropin inducible transcription repressor-3 /DB_XREF=gi:7706464 /UG=Hs.102397 GIOT-3 for gonadotropin inducible transcription repressor-3 /FL=gb:AB021643.1 gb:NM_016265.1	NM_016265		NP_057349
219575_s_at	0.010205	gb:NM_022341.1 /DEF=Homo sapiens peptide deformylase-like protein (LOC64146), mRNA. /FEA=mRNA /GEN=LOC64146 /PROD=peptide deformylase-like protein /DB_XREF=gi:11641242 /UG=Hs.130849 peptide deformylase-like protein /FL=gb:AF239156.1 gb:NM_022341.1 gb:AF322879.1	NM_022341		NP_071736

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219594_at	1.72E-04	gb:NM_016533.1 /DEF=Homo sapiens ninjurin 2 (NINJ2), mRNA. /FEA=mRNA /GEN=NINJ2 /PROD=ninjurin 2 /DB_XREF=gi:7706622 /UG=Hs.239208 ninjurin 2 /FL=gb:AF205633.1 gb:NM_016533.1	NM_016533		
219607_s_at	3.56E-04	gb:NM_024021.1 /DEF=Homo sapiens membrane-spanning 4-domains, subfamily A, member 4 (MS4A4), mRNA. /FEA=mRNA /GEN=MS4A4 /PROD=membrane-spanning 4-domains, subfamily A, member4 /DB_XREF=gi:13430865 /UG=Hs.325960 membrane-spanning 4-domains, subfamily A, member 4 /FL=gb:AB013102.1 gb:NM_024021.1 gb:AF068288.1 gb:NM_016650.1	NM_024021		NP_683876
219608_s_at	0.043703	gb:NM_024862.1 /DEF=Homo sapiens hypothetical protein FLJ13962 (FLJ13962), mRNA. /FEA=mRNA /GEN=FLJ13962 /PROD=hypothetical protein FLJ13962 /DB_XREF=gi:13376291 /UG=Hs.330407 hypothetical protein FLJ13962 /FL=gb:NM_024862.1	NM_024862		NP_079138
219620_x_at	5.03E-04	gb:NM_017723.1 /DEF=Homo sapiens hypothetical protein FLJ20245 (FLJ20245), mRNA. /FEA=mRNA /GEN=FLJ20245 /PROD=hypothetical protein FLJ20245 /DB_XREF=gi:8923220 /UG=Hs.169758 hypothetical protein FLJ20245 /FL=gb:NM_017723.1	NM_017723		NP_060193

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219625_s_at	0.003597	gb:NM_005713.1 /DEF=Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) binding protein (COL4A3BP), mRNA. /FEA=mRNA /GEN=COL4A3BP /PROD=collagen, type IV, alpha 3 (Goodpasture antigen)binding protein /DB_XREF=gi:5031716 /UG=Hs.21276 collagen, type IV, alpha 3 (Goodpasture antigen) binding protein /FL=gb:AF136450.1 gb:NM_005713.1	NM_005713		NP_112729
219632_s_at	0.005121	gb:AF196175.1 /DEF=Homo sapiens capsaicin receptor mRNA, complete cds. /FEA=mRNA /PROD=capsaicin receptor /DB_XREF=gi:12003145 /UG=Hs.283010 vanilloid receptor subtype 1 /FL=gb:AF196175.1 gb:NM_018727.1	AF196175		NP_542437
219634_at	0.010205	gb:NM_018413.1 /DEF=Homo sapiens chondroitin 4-sulfotransferase (C4ST), mRNA. /FEA=mRNA /GEN=C4ST /PROD=chondroitin 4-sulfotransferase /DB_XREF=gi:8923757 /UG=Hs.287402 chondroitin 4-sulfotransferase /FL=gb:AB042326.1 gb:NM_018413.1 gb:AF239820.1	NM_018413		NP_060883
219639_x_at	0.004341	gb:NM_020213.1 /DEF=Homo sapiens hypothetical protein from EUROIMAGE 1977056 (LOC56965), mRNA. /FEA=mRNA /GEN=LOC56965 /PROD=hypothetical protein from EUROIMAGE 1977056 /DB_XREF=gi:9910373 /UG=Hs.315687 hypothetical protein from EUROIMAGE 1977056 /FL=gb:NM_020213.1	NM_020213		NP_064599

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219644_at	0.013088	gb:NM_016122.1 /DEF=Homo sapiens NY-REN-58 antigen (LOC51134), mRNA. /FEA=mRNA /GEN=LOC51134 /PROD=NY-REN-58 antigen /DB_XREF=gi:7705838 /UG=Hs.56148 NY-REN-58 antigen /FL=gb:AF155115.1 gb:NM_016122.1	NM_016122		NP_057206
219649_at	3.26E-04	gb:NM_013339.1 /DEF=Homo sapiens dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase (ALG6), mRNA. /FEA=mRNA /GEN=ALG6 /PROD=dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase /DB_XREF=gi:7019324 /UG=Hs.80042 dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase /FL=gb:AF063604.1 gb:BC001253.1 gb:AF102851.1 gb:NM_013339.1	NM_013339		NP_037471
219657_s_at	0.013223	gb:NM_016531.1 /DEF=Homo sapiens hBKLF for basic kruppel like factor (LOC51274), mRNA. /FEA=mRNA /GEN=LOC51274 /PROD=hBKLF for basic kruppel like factor /DB_XREF=gi:7706076 /UG=Hs.119640 hBKLF for basic kruppel like factor /FL=gb:AF285837.1 gb:AB024523.1 gb:NM_016531.1	NM_016531		NP_057615
219666_at	1.26E-04	gb:NM_022349.1 /DEF=Homo sapiens CD20-like precursor (LOC64166), mRNA. /FEA=mRNA /GEN=LOC64166 /PROD=CD20-like precursor /DB_XREF=gi:11641258 /UG=Hs.17914 membrane-spanning 4-domains, subfamily A, member6 /FL=gb:AF142409.1 gb:NM_022349.1	NM_022349		NP_690591

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219667_s_at	0.007052	gb:NM_017935.1 /DEF=Homo sapiens hypothetical protein FLJ20706 (FLJ20706), mRNA. /FEA=mRNA /GEN=FLJ20706 /PROD=hypothetical protein FLJ20706 /DB_XREF=gi:8923635 /UG=Hs.193736 hypothetical protein FLJ20706 /FL=gb:NM_017935.1	NM_017935		NP_060405
219669_at	0.002536	gb:NM_020406.1 /DEF=Homo sapiens polycythemia rubra vera 1; cell surface receptor (PRV1), mRNA. /FEA=mRNA /GEN=PRV1 /PROD=polycythemia rubra vera 1; cell surfacereceptor /DB_XREF=gi:9966888 /UG=Hs.232165 polycythemia rubra vera 1; cell surface receptor /FL=gb:AF146747.1 gb:NM_020406.1	NM_020406		NP_065139
219672_at	0.008237	gb:NM_016633.1 /DEF=Homo sapiens EDRF protein (LOC51327), mRNA. /FEA=mRNA /GEN=LOC51327 /PROD=EDRF protein /DB_XREF=gi:7706179 /UG=Hs.274309 erythroid differentiation-related factor /FL=gb:AF208865.1 gb:NM_016633.1	NM_016633		NP_057717
219673_at	0.004341	gb:NM_017696.1 /DEF=Homo sapiens hypothetical protein FLJ20170 (FLJ20170), mRNA. /FEA=mRNA /GEN=FLJ20170 /PROD=hypothetical protein FLJ20170 /DB_XREF=gi:8923164 /UG=Hs.279008 hypothetical protein FLJ20170 /FL=gb:NM_017696.1	NM_017696		NP_060166
219680_at	0.041795	gb:NM_024618.1 /DEF=Homo sapiens hypothetical protein FLJ21478 (FLJ21478), mRNA. /FEA=mRNA /GEN=FLJ21478 /PROD=hypothetical protein FLJ21478 /DB_XREF=gi:13375837 /UG=Hs.31097 hypothetical protein FLJ21478 /FL=gb:NM_024618.1	NM_024618		NP_733840

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219694_at	1.26E-04	gb:NM_019018.1 /DEF=Homo sapiens hypothetical protein (FLJ11127), mRNA. /FEA=mRNA /GEN=FLJ11127 /PROD=hypothetical protein /DB_XREF=gi:9506640 /UG=Hs.91165 hypothetical protein /FL=gb:NM_019018.1	NM_019018		NP_061891
219700_at	3.56E-04	gb:NM_020405.1 /DEF=Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA. /FEA=mRNA /GEN=TEM7 /PROD=tumor endothelial marker 7 precursor /DB_XREF=gi:9966886 /UG=Hs.125036 tumor endothelial marker 7 precursor /FL=gb:AF279144.1 gb:NM_020405.1	NM_020405		NP_065138
219711_at	0.026891	gb:NM_017652.1 /DEF=Homo sapiens hypothetical protein FLJ20070 (FLJ20070), mRNA. /FEA=mRNA /GEN=FLJ20070 /PROD=hypothetical protein FLJ20070 /DB_XREF=gi:8923076 /UG=Hs.186711 hypothetical protein FLJ20070 /FL=gb:NM_017652.1	NM_017652		NP_060122
219717_at	6.68E-04	gb:NM_017741.1 /DEF=Homo sapiens hypothetical protein FLJ20280 (FLJ20280), mRNA. /FEA=mRNA /GEN=FLJ20280 /PROD=hypothetical protein FLJ20280 /DB_XREF=gi:8923256 /UG=Hs.270134 hypothetical protein FLJ20280 /FL=gb:NM_017741.1	NM_017741		NP_060211
219734_at	0.00161	gb:NM_017699.1 /DEF=Homo sapiens hypothetical protein FLJ20174 (FLJ20174), mRNA. /FEA=mRNA /GEN=FLJ20174 /PROD=hypothetical protein FLJ20174 /DB_XREF=gi:8923170 /UG=Hs.114556 hypothetical protein FLJ20174 /FL=gb:NM_017699.1	NM_017699		NP_060169

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219770_at	0.005852	gb:NM_024659.1 /DEF=Homo sapiens hypothetical protein FLJ11753 (FLJ11753), mRNA. /FEA=mRNA /GEN=FLJ11753 /PROD=hypothetical protein FLJ11753 /DB_XREF=gi:13375910 /UG=Hs.62348 hypothetical protein FLJ11753 /FL=gb:NM_024659.1	NM_024659		NP_078935
219797_at	7.58E-04	gb:NM_012214.1 /DEF=Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A (MGAT4A), mRNA. /FEA=mRNA /GEN=MGAT4A /PROD=mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A /DB_XREF=gi:6912501 /UG=Hs.177576 mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A /FL=gb:AB000616.1 gb:NM_012214.1	NM_012214		NP_036346
219799_s_at	0.033381	gb:NM_005771.1 /DEF=Homo sapiens retinol dehydrogenase homolog (RDHL), mRNA. /FEA=mRNA /GEN=RDHL /PROD=retinol dehydrogenase homolog /DB_XREF=gi:5032034 /UG=Hs.179608 retinol dehydrogenase homolog /FL=gb:AF067174.1 gb:NM_005771.1	NM_005771		NP_005762
219806_s_at	1.26E-04	gb:NM_020179.1 /DEF=Homo sapiens FN5 protein (FN5), mRNA. /FEA=mRNA /GEN=FN5 /PROD=FN5 protein /DB_XREF=gi:9910225 /UG=Hs.259737 FN5 protein /FL=gb:AF197137.1 gb:NM_020179.1	NM_020179		NP_064564

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219812_at	6.68E-04	gb:NM_024070.1 /DEF=Homo sapiens hypothetical protein MGC2463 (MGC2463), mRNA. /FEA=mRNA /GEN=MGC2463 /PROD=hypothetical protein MGC2463 /DB_XREF=gi:13129051 /UG=Hs.323634 hypothetical protein MGC2463 /FL=gb:BC001129.1 gb:NM_024070.1	NM_024070		
219816_s_at	3.26E-04	gb:NM_018107.1 /DEF=Homo sapiens hypothetical protein FLJ10482 (FLJ10482), mRNA. /FEA=mRNA /GEN=FLJ10482 /PROD=hypothetical protein FLJ10482 /DB_XREF=gi:8922449 /UG=Hs.4997 hypothetical protein FLJ10482 /FL=gb:NM_018107.1	NM_018107		NP_060577
219846_at	0.016934	gb:NM_025174.1 /DEF=Homo sapiens hypothetical protein FLJ23040 (FLJ23040), mRNA. /FEA=mRNA /GEN=FLJ23040 /PROD=hypothetical protein FLJ23040 /DB_XREF=gi:13376769 /UG=Hs.169813 hypothetical protein FLJ23040 /FL=gb:NM_025174.1	NM_025174		NP_079450
219859_at	1.26E-04	gb:NM_014358.1 /DEF=Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9 (CLECSF9), mRNA. /FEA=mRNA /GEN=CLECSF9 /PROD=C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9 /DB_XREF=gi:7657332 /UG=Hs.236516 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9 /FL=gb:BC000715.1 gb:AB024718.1 gb:NM_014358.1	NM_014358		NP_055173

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219860_at	0.010205	gb:NM_025262.1 /DEF=Homo sapiens G5C protein (G5C), mRNA. /FEA=mRNA /GEN=G5C /PROD=G5C protein /DB_XREF=gi:13376875 /UG=Hs.246845 G5C protein /FL=gb:NM_025262.1	NM_025262		NP_079538
219862_s_at	0.007779	gb:NM_012336.1 /DEF=Homo sapiens nuclear prelamin A recognition factor (NARF), mRNA. /FEA=mRNA /GEN=NARF /PROD=nuclear prelamin A recognition factor /DB_XREF=gi:6912523 /UG=Hs.256526 nuclear prelamin A recognition factor /FL=gb:AF128406.1 gb:NM_012336.1	NM_012336		NP_114174
219868_s_at	0.039278	gb:NM_016376.1 /DEF=Homo sapiens ANKHZN protein (ANKHZN), mRNA. /FEA=mRNA /GEN=ANKHZN /PROD=ANKHZN protein /DB_XREF=gi:7705277 /UG=Hs.284163 ANKHZN protein /FL=gb:AB037360.1 gb:NM_016376.1	NM_016376		NP_065791
219869_s_at	9.79E-04	gb:NM_022154.1 /DEF=Homo sapiens up-regulated by BCG-CWS (LOC64116), mRNA. /FEA=mRNA /GEN=LOC64116 /PROD=up-regulated by BCG-CWS /DB_XREF=gi:11545899 /UG=Hs.284205 up-regulated by BCG-CWS /FL=gb:NM_022154.1	NM_022154		NP_071437
219885_at	0.046329	gb:NM_018042.1 /DEF=Homo sapiens hypothetical protein FLJ10260 (FLJ10260), mRNA. /FEA=mRNA /GEN=FLJ10260 /PROD=hypothetical protein FLJ10260 /DB_XREF=gi:8922317 /UG=Hs.105859 hypothetical protein FLJ10260 /FL=gb:NM_018042.1	NM_018042		NP_060512

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219892_at	1.26E-04	gb:NM_023003.1 /DEF=Homo sapiens transmembrane 6 superfamily member 1 (TM6SF1), mRNA. /FEA=mRNA /GEN=TM6SF1 /PROD=transmembrane 6 superfamily member 1 /DB_XREF=gi:13194198 /UG=Hs.133865 transmembrane 6 superfamily member 1 /FL=gb:AF255922.1 gb:NM_023003.1	NM_023003		NP_075379
219929_s_at	8.34E-04	gb:NM_024071.1 /DEF=Homo sapiens hypothetical protein MGC2550 (MGC2550), mRNA. /FEA=mRNA /GEN=MGC2550 /PROD=hypothetical protein MGC2550 /DB_XREF=gi:13129053 /UG=Hs.318498 hypothetical protein MGC2550 /FL=gb:BC001130.1 gb:NM_024071.1	NM_024071		NP_076976
219931_s_at	0.002536	gb:NM_021633.1 /DEF=Homo sapiens kelch-like protein C3IP1 (C3IP1), mRNA. /FEA=mRNA /GEN=C3IP1 /PROD=kelch-like protein C3IP1 /DB_XREF=gi:11056005 /UG=Hs.3826 kelch-like protein C3IP1 /FL=gb:AF190900.1 gb:NM_021633.1 gb:BC003183.1 gb:BC004175.1	NM_021633		NP_067646
219933_at	0.016934	gb:NM_016066.1 /DEF=Homo sapiens CGI-133 protein (LOC51022), mRNA. /FEA=mRNA /GEN=LOC51022 /PROD=CGI-133 protein /DB_XREF=gi:7705627 /UG=Hs.5054 CGI-133 protein /FL=gb:AF151891.1 gb:NM_016066.1 gb:AF132495.2	NM_016066		NP_057150

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219938_s_at	3.26E-04	gb:NM_024430.1 /DEF=Homo sapiens proline-serine-threonine phosphatase interacting protein 2 (PSTPIP2), mRNA. /FEA=mRNA /GEN=PSTPIP2 /PROD=proline-serine-threonine phosphatase interactingprotein 2 /DB_XREF=gi:13270468 /UG=Hs.69149 proline-serine-threonine phosphatase interacting protein 2 /FL=gb:NM_024430.1	NM_024430		NP_077748
219947_at	0.001116	gb:NM_016184.1 /DEF=Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 (CLECSF6), mRNA. /FEA=mRNA /GEN=CLECSF6 /PROD=C-type (calcium dependent,carbohydrate-recognition domain) lectin, superfamily member 6 /DB_XREF=gi:7705337 /UG=Hs.115515 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /FL=gb:AF109146.1 gb:NM_016184.1 gb:AF067800.1	NM_016184		NP_057268
219952_s_at	0.007934	gb:NM_020533.1 /DEF=Homo sapiens mucolipin 1 (MCOLN1), mRNA. /FEA=mRNA /GEN=MCOLN1 /PROD=mucolipin 1 /DB_XREF=gi:10092596 /UG=Hs.12909 mucolipin 1 /FL=gb:NM_020533.1 gb:BC005149.1 gb:AF287269.1 gb:AF249319.1	NM_020533		NP_065394

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219960_s_at	0.010205	gb:NM_015984.1 /DEF=Homo sapiens ubiquitin C-terminal hydrolase UCH37 (UCH37), mRNA. /FEA=mRNA /GEN=UCH37 /PROD=ubiquitin C-terminal hydrolase UCH37 /DB_XREF=gi:7706752 /UG=Hs.171581 ubiquitin C-terminal hydrolase UCH37 /FL=gb:AF147717.1 gb:NM_015984.1	NM_015984		NP_057068
219963_at	0.004659	gb:NM_016364.1 /DEF=Homo sapiens protein phosphatase (LOC51207), mRNA. /FEA=mRNA /GEN=LOC51207 /PROD=protein phosphatase /DB_XREF=gi:7705958 /UG=Hs.178170 protein phosphatase /FL=gb:AB027004.1 gb:NM_016364.1	NM_016364		NP_057448
219974_x_at	0.02145	gb:NM_018479.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HCDASE (LOC55862), mRNA. /FEA=mRNA /GEN=LOC55862 /PROD=uncharacterized hypothalamus protein HCDASE /DB_XREF=gi:8923863 /UG=Hs.239218 uncharacterized hypothalamus protein HCDASE /FL=gb:AF220192.1 gb:NM_018479.1	NM_018479		NP_060949
219979_s_at	0.003171	gb:NM_016401.1 /DEF=Homo sapiens hypothetical protein (HSPC138), mRNA. /FEA=mRNA /GEN=HSPC138 /PROD=hypothetical protein /DB_XREF=gi:7705470 /UG=Hs.283775 hypothetical protein /FL=gb:AF161487.1 gb:NM_016401.1	NM_016401		NP_057485
219981_x_at	0.001116	gb:NM_017961.1 /DEF=Homo sapiens hypothetical protein FLJ20813 (FLJ20813), mRNA. /FEA=mRNA /GEN=FLJ20813 /PROD=hypothetical protein FLJ20813 /DB_XREF=gi:8923685 /UG=Hs.306203 hypothetical protein FLJ20813 /FL=gb:NM_017961.1	NM_017961		NP_060431

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219994_at	0.028893	gb:NM_019043.1 /DEF=Homo sapiens similar to proline-rich protein 48 (LOC54518), mRNA. /FEA=mRNA /GEN=LOC54518 /PROD=similar to proline-rich protein 48 /DB_XREF=gi:9506856 /UG=Hs.98874 similar to proline-rich protein 48 /FL=gb:NM_019043.1	NM_019043		NP_061916
220000_at	3.09E-04	gb:NM_003830.1 /DEF=Homo sapiens sialic acid binding Ig-like lectin 5 (SIGLEC5), mRNA. /FEA=mRNA /GEN=SIGLEC5 /PROD=sialic acid binding Ig-like lectin 5 /DB_XREF=gi:4502658 /UG=Hs.117005 sialic acid binding Ig-like lectin 5 /FL=gb:U71383.1 gb:NM_003830.1 gb:AF170484.1	NM_003830		NP_003821
220001_at	0.003171	gb:NM_012387.1 /DEF=Homo sapiens peptidyl arginine deiminase, type V (PAD), mRNA. /FEA=mRNA /GEN=PAD /PROD=peptidyl arginine deiminase, type V /DB_XREF=gi:6912575 /UG=Hs.117232 peptidyl arginine deiminase, type V /FL=gb:AB017919.1 gb:NM_012387.1	NM_012387		NP_036519
220005_at	1.26E-04	gb:NM_023914.1 /DEF=Homo sapiens G protein-coupled receptor 86 (GPR86), mRNA. /FEA=mRNA /GEN=GPR86 /PROD=G protein-coupled receptor 86 /DB_XREF=gi:13194202 /UG=Hs.13040 G protein-coupled receptor 86 /FL=gb:AF295368.1 gb:NM_023914.1 gb:AF178982.1 gb:AF345565.1	NM_023914		NP_795713

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220009_at	0.008237	gb:NM_024778.1 /DEF=Homo sapiens hypothetical protein FLJ22612 (FLJ22612), mRNA. /FEA=mRNA /GEN=FLJ22612 /PROD=hypothetical protein FLJ22612 /DB_XREF=gi:13376133 /UG=Hs.144266 hypothetical protein FLJ22612 /FL=gb:NM_024778.1	NM_024778		NP_079054
220015_at	0.004341	gb:NM_017766.1 /DEF=Homo sapiens hypothetical protein FLJ20321 (FLJ20321), mRNA. /FEA=mRNA /GEN=FLJ20321 /PROD=hypothetical protein FLJ20321 /DB_XREF=gi:8923302 /UG=Hs.162196 hypothetical protein FLJ20321 /FL=gb:BC004410.1 gb:NM_017766.1	NM_017766		NP_060236
220023_at	0.016934	gb:NM_018690.1 /DEF=Homo sapiens apolipoprotein B48 receptor (APOB48R), mRNA. /FEA=mRNA /GEN=APOB48R /PROD=apolipoprotein B48 receptor /DB_XREF=gi:8922078 /UG=Hs.200333 apolipoprotein B48 receptor /FL=gb:AF141332.1 gb:AF141334.1 gb:NM_018690.1	NM_018690		NP_061160
220034_at	0.003522	gb:NM_007199.1 /DEF=Homo sapiens interleukin-1 receptor-associated kinase M (IRAK-M), mRNA. /FEA=mRNA /GEN=IRAK-M /PROD=interleukin-1 receptor-associated kinase M /DB_XREF=gi:6005791 /UG=Hs.268552 interleukin-1 receptor-associated kinase M /FL=gb:AF113136.1 gb:NM_007199.1	NM_007199		NP_009130

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220050_at	0.00423	gb:NM_018956.1 /DEF=Homo sapiens chromosome 9 open reading frame 9 (C9ORF9), mRNA. /FEA=mRNA /GEN=C9ORF9 /PROD=chromosome 9 open reading frame 9 /DB_XREF=gi:9506440 /UG=Hs.62595 chromosome 9 open reading frame 9 /FL=gb:NM_018956.1	NM_018956		NP_061829
220052_s_at	0.00161	gb:NM_012461.1 /DEF=Homo sapiens TERF1 (TRF1)-interacting nuclear factor 2 (TINF2), mRNA. /FEA=mRNA /GEN=TINF2 /PROD=TERF1 (TRF1)-interacting nuclear factor 2 /DB_XREF=gi:6912715 /UG=Hs.7797 TERF1 (TRF1)-interacting nuclear factor 2 /FL=gb:AF195512.1 gb:NM_012461.1	NM_012461		NP_036593
220071_x_at	0.00228	gb:NM_018097.1 /DEF=Homo sapiens hypothetical protein FLJ10460 (FLJ10460), mRNA. /FEA=mRNA /GEN=FLJ10460 /PROD=hypothetical protein FLJ10460 /DB_XREF=gi:8922429 /UG=Hs.14347 hypothetical protein FLJ10460 /FL=gb:NM_018097.1	NM_018097		NP_060567
220078_at	0.001618	gb:NM_018391.1 /DEF=Homo sapiens hypothetical protein FLJ11328 (FLJ11328), mRNA. /FEA=mRNA /GEN=FLJ11328 /PROD=hypothetical protein FLJ11328 /DB_XREF=gi:8922996 /UG=Hs.178518 hypothetical protein FLJ11328 /FL=gb:NM_018391.1	NM_018391		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220088_at	0.001116	gb:NM_001736.1 /DEF=Homo sapiens complement component 5 receptor 1 (C5a ligand) (C5R1), mRNA. /FEA=mRNA /GEN=C5R1 /PROD=complement component 5 receptor 1 (C5a ligand) /DB_XREF=gi:4502508 /UG=Hs.2161 complement component 5 receptor 1 (C5a ligand) /FL=gb:M62505.1 gb:NM_001736.1	NM_001736		NP_001727
220097_s_at	0.046435	gb:NM_017728.1 /DEF=Homo sapiens hypothetical protein FLJ20255 (FLJ20255), mRNA. /FEA=mRNA /GEN=FLJ20255 /PROD=hypothetical protein FLJ20255 /DB_XREF=gi:8923229 /UG=Hs.274247 hypothetical protein FLJ20255 /FL=gb:NM_017728.1	NM_017728		NP_060198
220112_at	0.00161	gb:NM_024669.1 /DEF=Homo sapiens hypothetical protein FLJ11795 (FLJ11795), mRNA. /FEA=mRNA /GEN=FLJ11795 /PROD=hypothetical protein FLJ11795 /DB_XREF=gi:13375927 /UG=Hs.84560 hypothetical protein FLJ11795 /FL=gb:NM_024669.1	NM_024669		NP_078945
220113_x_at	0.001116	gb:NM_019014.1 /DEF=Homo sapiens similar to DNA-directed RNA polymerase I (135 kDa) (FLJ10816), mRNA. /FEA=mRNA /GEN=FLJ10816 /PROD=similar to DNA-directed RNA polymerase I (135kDa) /DB_XREF=gi:9506618 /UG=Hs.86337 similar to DNA-directed RNA polymerase I (135 kDa) /FL=gb:NM_019014.1	NM_019014		NP_115588

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220122_at	0.033381	gb:NM_024717.1 /DEF=Homo sapiens hypothetical protein FLJ22344 (FLJ22344), mRNA. /FEA=mRNA /GEN=FLJ22344 /PROD=hypothetical protein FLJ22344 /DB_XREF=gi:13376020 /UG=Hs.107716 hypothetical protein FLJ22344 /FL=gb:NM_024717.1	NM_024717		NP_078993
220127_s_at	0.041055	gb:NM_017703.1 /DEF=Homo sapiens hypothetical protein FLJ20188 (FLJ20188), mRNA. /FEA=mRNA /GEN=FLJ20188 /PROD=hypothetical protein FLJ20188 /DB_XREF=gi:8923178 /UG=Hs.12439 hypothetical protein FLJ20188 /FL=gb:NM_017703.1	NM_017703		NP_060173
220132_s_at	0.016934	gb:NM_013269.1 /DEF=Homo sapiens lectin-like NK cell receptor (LLT1), mRNA. /FEA=mRNA /GEN=LLT1 /PROD=lectin-like NK cell receptor /DB_XREF=gi:7019446 /UG=Hs.136748 lectin-like NK cell receptor /FL=gb:AF133299.1 gb:NM_013269.1 gb:AF285087.1	NM_013269		NP_037401
220138_at	5.07E-04	gb:NM_004821.1 /DEF=Homo sapiens heart and neural crest derivatives expressed 1 (HAND1), mRNA. /FEA=mRNA /GEN=HAND1 /PROD=basic helix-loop-helix transcription factorHAND1 /DB_XREF=gi:4758505 /UG=Hs.152531 heart and neural crest derivatives expressed 1 /FL=gb:AF061756.1 gb:NM_004821.1	NM_004821		NP_004812
220140_s_at	0.033381	gb:NM_013323.1 /DEF=Homo sapiens sorting nexin 11 (SNX11), mRNA. /FEA=mRNA /GEN=SNX11 /PROD=sorting nexin 11 /DB_XREF=gi:7019538 /UG=Hs.15827 sorting nexin 11 /FL=gb:AF121861.1 gb:NM_013323.1	NM_013323		NP_689450

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220147_s_at	0.013223	gb:NM_021238.1 /DEF=Homo sapiens TERA protein (TERA), mRNA. /FEA=mRNA /GEN=TERA /PROD=TERA protein /DB_XREF=gi:10864048 /UG=Hs.180780 TERA protein /FL=gb:NM_021238.1 gb:BC000024.1	NM_021238		NP_067061
220162_s_at	0.001012	gb:NM_022352.1 /DEF=Homo sapiens caspase recruitment domain protein 9 (LOC64170), mRNA. /FEA=mRNA /GEN=LOC64170 /PROD=caspase recruitment domain protein 9 /DB_XREF=gi:11641262 /UG=Hs.271815 caspase recruitment domain protein 9 /FL=gb:NM_022352.1	NM_022352		NP_434701
220175_s_at	0.041055	gb:NM_020667.1 /DEF=Homo sapiens hypothetical protein from clone 1659351 (LOC57397), mRNA. /FEA=mRNA /GEN=LOC57397 /PROD=hypothetical protein from clone 1659351 /DB_XREF=gi:10190707 /UG=Hs.288838 hypothetical protein from clone 1659351 /FL=gb:NM_020667.1	NM_020667		
220202_s_at	0.00228	gb:NM_018835.1 /DEF=Homo sapiens membrane-associated nucleic acid binding protein (MNAB), mRNA. /FEA=mRNA /GEN=MNAB /PROD=membrane-associated nucleic acid bindingprotein /DB_XREF=gi:9256536 /UG=Hs.112227 membrane-associated nucleic acid binding protein /FL=gb:NM_018835.1	NM_018835		NP_061323

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220232_at	0.042536	gb:NM_024906.1 /DEF=Homo sapiens hypothetical protein FLJ21032 (FLJ21032), mRNA. /FEA=mRNA /GEN=FLJ21032 /PROD=hypothetical protein FLJ21032 /DB_XREF=gi:13376362 /UG=Hs.247474 hypothetical protein FLJ21032 /FL=gb:NM_024906.1	NM_024906		NP_079182
220235_s_at	1.26E-04	gb:NM_018372.1 /DEF=Homo sapiens hypothetical protein FLJ11269 (FLJ11269), mRNA. /FEA=mRNA /GEN=FLJ11269 /PROD=hypothetical protein FLJ11269 /DB_XREF=gi:8922961 /UG=Hs.25245 hypothetical protein FLJ11269 /FL=gb:NM_018372.1	NM_018372		NP_060842
220244_at	0.007779	gb:NM_013343.1 /DEF=Homo sapiens NAG-7 protein (NAG-7), mRNA. /FEA=mRNA /GEN=NAG-7 /PROD=NAG-7 protein /DB_XREF=gi:7106370 /UG=Hs.278951 NAG-7 protein /FL=gb:AF086709.2 gb:NM_013343.1	NM_013343		NP_037475
220292_at	0.002464	gb:NM_024340.1 /DEF=Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA. /FEA=mRNA /GEN=MGC4179 /PROD=hypothetical protein MGC4179 /DB_XREF=gi:13236590 /UG=Hs.129369 hypothetical protein MGC4179 /FL=gb:BC002859.1 gb:NM_024340.1	NM_024340		
220305_at	3.26E-04	gb:NM_024030.1 /DEF=Homo sapiens hypothetical protein MGC3260 (MGC3260), mRNA. /FEA=mRNA /GEN=MGC3260 /PROD=hypothetical protein MGC3260 /DB_XREF=gi:13128977 /UG=Hs.15514 hypothetical protein MGC3260 /FL=gb:BC000073.1 gb:NM_024030.1	NM_024030		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220306_at	0.004026	gb:NM_017709.1 /DEF=Homo sapiens hypothetical protein FLJ20202 (FLJ20202), mRNA. /FEA=mRNA /GEN=FLJ20202 /PROD=hypothetical protein FLJ20202 /DB_XREF=gi:8923191 /UG=Hs.155556 hypothetical protein FLJ20202 /FL=gb:NM_017709.1	NM_017709		NP_060179
220330_s_at	1.26E-04	gb:NM_022136.1 /DEF=Homo sapiens SAM domain, SH3 domain and nuclear localisation signals, 1 (SAMSN1), mRNA. /FEA=mRNA /GEN=SAMSN1 /PROD=SAM domain, SH3 domain and nuclear localisationsignals, 1 /DB_XREF=gi:11545870 /UG=Hs.24633 SAM domain, SH3 domain and nuclear localisation signals, 1 /FL=gb:AF222927.1 gb:NM_022136.1	NM_022136		NP_071419
220363_s_at	0.004452	gb:NM_022086.1 /DEF=Homo sapiens hypothetical protein FLJ11656 (FLJ11656), mRNA. /FEA=mRNA /GEN=FLJ11656 /PROD=hypothetical protein FLJ11656 /DB_XREF=gi:11545798 /UG=Hs.96560 hypothetical protein FLJ11656 /FL=gb:NM_022086.1	NM_022086		NP_573403
220367_s_at	2.05E-04	gb:NM_024545.1 /DEF=Homo sapiens hypothetical protein FLJ12761 (FLJ12761), mRNA. /FEA=mRNA /GEN=FLJ12761 /PROD=hypothetical protein FLJ12761 /DB_XREF=gi:13375706 /UG=Hs.10554 hypothetical protein FLJ12761 /FL=gb:NM_024545.1	NM_024545		NP_078821

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220368_s_at	0.00228	gb:NM_017936.1 /DEF=Homo sapiens hypothetical protein FLJ20707 (FLJ20707), mRNA. /FEA=mRNA /GEN=FLJ20707 /PROD=hypothetical protein FLJ20707 /DB_XREF=gi:8923637 /UG=Hs.109441 hypothetical protein FLJ20707 /FL=gb:NM_017936.1	NM_017936		NP_115949
220375_s_at	0.035763	gb:NM_024752.1 /DEF=Homo sapiens hypothetical protein FLJ23312 (FLJ23312), mRNA. /FEA=mRNA /GEN=FLJ23312 /PROD=hypothetical protein FLJ23312 /DB_XREF=gi:13376081 /UG=Hs.126555 hypothetical protein FLJ23312 /FL=gb:NM_024752.1	NM_024752		
220404_at	0.010205	gb:NM_014076.1 /DEF=Homo sapiens PRO0611 protein (PRO0611), mRNA. /FEA=mRNA /GEN=PRO0611 /PROD=PRO0611 protein /DB_XREF=gi:7662571 /UG=Hs.163833 PRO0611 protein /FL=gb:AF111851.1 gb:NM_014076.1	NM_014076		
220416_at	1.26E-04	gb:NM_024837.1 /DEF=Homo sapiens hypothetical protein FLJ21472 (FLJ21472), mRNA. /FEA=mRNA /GEN=FLJ21472 /PROD=hypothetical protein FLJ21472 /DB_XREF=gi:13376248 /UG=Hs.182738 hypothetical protein FLJ21472 /FL=gb:NM_024837.1	NM_024837		NP_079113
220417_s_at	0.013223	gb:NM_015963.1 /DEF=Homo sapiens CGI-36 protein (LOC51078), mRNA. /FEA=mRNA /GEN=LOC51078 /PROD=CGI-36 protein /DB_XREF=gi:7705731 /UG=Hs.182986 CGI-36 protein /FL=gb:AF132970.1 gb:NM_015963.1	NM_015963		NP_057047

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220477_s_at	0.034221	gb:NM_014145.1 /DEF=Homo sapiens HSPC274 protein (HSPC274), mRNA. /FEA=mRNA /GEN=HSPC274 /PROD=HSPC274 protein /DB_XREF=gi:7661833 /UG=Hs.323476 HSPC274 protein /FL=gb:AF161392.1 gb:NM_014145.1	NM_014145		NP_054864
220491_at	0.014208	gb:NM_021175.1 /DEF=Homo sapiens liver-expressed antimicrobial peptide (LEAP-1), mRNA. /FEA=mRNA /GEN=LEAP-1 /PROD=liver-expressed antimicrobial peptide /DB_XREF=gi:10863972 /UG=Hs.8821 liver-expressed antimicrobial peptide /FL=gb:AF309489.1 gb:NM_021175.1 gb:AF131292.1	NM_021175		NP_066998
220494_s_at	0.043705	gb:NM_018678.1 /DEF=Homo sapiens lipopolysaccharide specific response-68 protein (LSR68), mRNA. /FEA=mRNA /GEN=LSR68 /PROD=lipopolysaccharide specific response-68 protein /DB_XREF=gi:8923914 /UG=Hs.103189 lipopolysaccharide specific response-68 protein /FL=gb:AF147723.1 gb:NM_018678.1	NM_018678		NP_061148
220500_s_at	0.009275	gb:NM_007082.2 /DEF=Homo sapiens RAB, member of RAS oncogene family-like 2A (RABL2A), mRNA. /FEA=mRNA /GEN=RABL2A /PROD=RAB, member of RAS oncogene family-like 2A /DB_XREF=gi:7549818 /UG=Hs.122660 RAB, member of RAS oncogene family-like 2A /FL=gb:AF095350.1 gb:NM_007082.2	NM_007082		NP_038198

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220532_s_at	0.013223	gb:NM_014020.1 /DEF=Homo sapiens LR8 protein (LR8), mRNA. /FEA=mRNA /GEN=LR8 /PROD=LR8 protein /DB_XREF=gi:7662497 /UG=Hs.190161 LR8 protein /FL=gb:AF115384.1 gb:NM_014020.1	NM_014020		NP_054739
220537_at	0.007063	gb:NM_017677.1 /DEF=Homo sapiens hypothetical protein FLJ20126 (FLJ20126), mRNA. /FEA=mRNA /GEN=FLJ20126 /PROD=hypothetical protein FLJ20126 /DB_XREF=gi:8923125 /UG=Hs.196568 hypothetical protein FLJ20126 /FL=gb:NM_017677.1	NM_017677		NP_060147
220570_at	0.041795	gb:NM_020415.2 /DEF=Homo sapiens found in inflammatory zone 3 (FIZZ3), mRNA. /FEA=mRNA /GEN=FIZZ3 /PROD=resistin /DB_XREF=gi:13435379 /UG=Hs.283091 found in inflammatory zone 3 /FL=gb:AF323081.1 gb:NM_020415.2 gb:AF205952.1	NM_020415		NP_065148
220577_at	1.26E-04	gb:NM_025006.1 /DEF=Homo sapiens hypothetical protein FLJ13373 (FLJ13373), mRNA. /FEA=mRNA /GEN=FLJ13373 /PROD=hypothetical protein FLJ13373 /DB_XREF=gi:13376522 /UG=Hs.287567 hypothetical protein FLJ13373 /FL=gb:NM_025006.1	NM_025006		NP_079282
220591_s_at	0.005852	gb:NM_025184.1 /DEF=Homo sapiens hypothetical protein FLJ22843 (FLJ22843), mRNA. /FEA=mRNA /GEN=FLJ22843 /PROD=hypothetical protein FLJ22843 /DB_XREF=gi:13376775 /UG=Hs.301143 hypothetical protein FLJ22843 /FL=gb:NM_025184.1	NM_025184		NP_079460

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220597_s_at	0.02145	gb:NM_018694.1 /DEF=Homo sapiens HSVI binding protein (LOC55913), mRNA. /FEA=mRNA /GEN=LOC55913 /PROD=HSV binding protein /DB_XREF=gi:8923906 /UG=Hs.306208 HSVI binding protein /FL=gb:AF267748.1 gb:NM_018694.1	NM_018694		NP_061164
220603_s_at	2.05E-04	gb:NM_018349.1 /DEF=Homo sapiens hypothetical protein FLJ11175 (FLJ11175), mRNA. /FEA=mRNA /GEN=FLJ11175 /PROD=hypothetical protein FLJ11175 /DB_XREF=gi:8922916 /UG=Hs.33368 hypothetical protein FLJ11175 /FL=gb:NM_018349.1	NM_018349		NP_060819
220610_s_at	0.029713	gb:NM_006309.1 /DEF=Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2), mRNA. /FEA=mRNA /GEN=LRRFIP2 /PROD=leucine rich repeat (in FLII) interacting protein 2 /DB_XREF=gi:5453725 /UG=Hs.57672 leucine rich repeat (in FLII) interacting protein 2 /FL=gb:AF115509.1 gb:NM_006309.1	NM_006309		NP_060194
220615_s_at	2.05E-04	gb:NM_018099.1 /DEF=Homo sapiens hypothetical protein FLJ10462 (FLJ10462), mRNA. /FEA=mRNA /GEN=FLJ10462 /PROD=hypothetical protein FLJ10462 /DB_XREF=gi:8922433 /UG=Hs.100895 hypothetical protein FLJ10462 /FL=gb:AL136843.1 gb:NM_018099.1	NM_018099		NP_060569
220661_s_at	1.26E-04	gb:NM_017865.1 /DEF=Homo sapiens hypothetical protein FLJ20531 (FLJ20531), mRNA. /FEA=mRNA /GEN=FLJ20531 /PROD=hypothetical protein FLJ20531 /DB_XREF=gi:8923497 /UG=Hs.23617 hypothetical protein FLJ20531 /FL=gb:NM_017865.1	NM_017865		NP_060335

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220669_at	0.041795	gb:NM_017493.2 /DEF=Homo sapiens Hin-1 (HSHIN1), mRNA. /FEA=mRNA /GEN=HSHIN1 /PROD=Hin-1 /DB_XREF=gi:13518029 /UG=Hs.252722 Hin-1 /FL=gb:NM_017493.2	NM_017493		NP_059963
220684_at	0.003171	gb:NM_013351.1 /DEF=Homo sapiens T-box 21 (TBX21), mRNA. /FEA=mRNA /GEN=TBX21 /PROD=T-box 21 /DB_XREF=gi:7019548 /UG=Hs.272409 T-box 21 /FL=gb:AF093098.1 gb:NM_013351.1 gb:AF241243.2	NM_013351		NP_037483
220691_at	0.026891	gb:NM_014114.1 /DEF=Homo sapiens PRO0097 protein (PRO0097), mRNA. /FEA=mRNA /GEN=PRO0097 /PROD=PRO0097 protein /DB_XREF=gi:7662521 /UG=Hs.278928 PRO0097 protein /FL=gb:AF090890.1 gb:NM_014114.1	NM_014114		NP_054833
220710_at	0.007258	gb:NM_024970.1 /DEF=Homo sapiens hypothetical protein FLJ11722 (FLJ11722), mRNA. /FEA=mRNA /GEN=FLJ11722 /PROD=hypothetical protein FLJ11722 /DB_XREF=gi:13376465 /UG=Hs.287444 hypothetical protein FLJ11722 /FL=gb:NM_024970.1	NM_024970		NP_079246
220739_s_at	1.26E-04	gb:NM_017623.1 /DEF=Homo sapiens hypothetical protein FLJ20018 (FLJ20018), mRNA. /FEA=mRNA /GEN=FLJ20018 /PROD=hypothetical protein FLJ20018 /DB_XREF=gi:8923022 /UG=Hs.44095 hypothetical protein FLJ20018 /FL=gb:NM_017623.1	NM_017623		NP_060093

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220741_s_at	1.26E-04	gb:NM_006903.1 /DEF=Homo sapiens inorganic pyrophosphatase (SID6-306), mRNA. /FEA=mRNA /GEN=SID6-306 /PROD=inorganic pyrophosphatase /DB_XREF=gi:13491177 /UG=Hs.5123 inorganic pyrophosphatase /FL=gb:NM_006903.1 gb:AB026722.1	NM_006903		NP_789845
220746_s_at	0.041055	gb:NM_016290.1 /DEF=Homo sapiens retinoid x receptor interacting protein (LOC51720), mRNA. /FEA=mRNA /GEN=LOC51720 /PROD=retinoid x receptor interacting protein /DB_XREF=gi:7706482 /UG=Hs.7889 retinoid x receptor interacting protein /FL=gb:AF113538.1 gb:NM_016290.1	NM_016290		NP_057374
220750_s_at	0.013223	gb:NM_022356.1 /DEF=Homo sapiens growth suppressor 1 (GROS1), mRNA. /FEA=mRNA /GEN=GROS1 /PROD=growth suppressor 1 /DB_XREF=gi:11641270 /UG=Hs.10114 growth suppressor 1 /FL=gb:AF097431.1 gb:NM_022356.1	NM_022356		NP_071751
220753_s_at	0.048741	gb:NM_015974.1 /DEF=Homo sapiens lambda-crystallin (LOC51084), mRNA. /FEA=mRNA /GEN=LOC51084 /PROD=lambda-crystallin /DB_XREF=gi:7705743 /UG=Hs.108896 lambda-crystallin /FL=gb:AF077049.1 gb:NM_015974.1	NM_015974		NP_057058

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220757_s_at	0.02145	gb:NM_025241.1 /DEF=Homo sapiens UBX domain-containing gene 1 (UBXD1), mRNA. /FEA=mRNA /GEN=UBXD1 /PROD=UBX domain-containing gene 1 /DB_XREF=gi:13376853 /UG=Hs.11081 UBX domain-containing 2 /FL=gb:AF272893.1 gb:NM_025241.1	NM_025241		NP_079517
220761_s_at	0.00161	gb:NM_016281.1 /DEF=Homo sapiens STE20-like kinase (JIK), mRNA. /FEA=mRNA /GEN=JIK /PROD=STE20-like kinase /DB_XREF=gi:7705559 /UG=Hs.12040 STE20-like kinase /FL=gb:AF179867.1 gb:NM_016281.1	NM_016281		NP_057365
220796_x_at	0.033381	gb:NM_024881.1 /DEF=Homo sapiens hypothetical protein FLJ14251 (FLJ14251), mRNA. /FEA=mRNA /GEN=FLJ14251 /PROD=hypothetical protein FLJ14251 /DB_XREF=gi:13376323 /UG=Hs.214178 hypothetical protein FLJ14251 /FL=gb:NM_024881.1	NM_024881		NP_079157
220800_s_at	0.018693	gb:NM_014547.1 /DEF=Homo sapiens tropomodulin 3 (ubiquitous) (TMOD3), mRNA. /FEA=mRNA /GEN=TMOD3 /PROD=tropomodulin 3 (ubiquitous) /DB_XREF=gi:7657648 /UG=Hs.22826 tropomodulin 3 (ubiquitous) /FL=gb:AF237631.1 gb:NM_014547.1	NM_014547		NP_055362
220864_s_at	0.005852	gb:NM_015965.1 /DEF=Homo sapiens CGI-39 protein; cell death-regulatory protein GRIM19 (LOC51079), mRNA. /FEA=mRNA /GEN=LOC51079 /PROD=CGI-39 protein; cell death-regulatory protein GRIM19 /DB_XREF=gi:7705733 /UG=Hs.279574 CGI-39 protein; cell death-regulatory protein GRIM19 /FL=gb:AF132973.1 gb:AF155662.1 gb:NM_015965.1	NM_015965		NP_057049

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220865_s_at	5.03E-04	gb:NM_014317.2 /DEF=Homo sapiens trans-prenyltransferase (TPT), mRNA. /FEA=mRNA /GEN=TPT /PROD=trans-prenyltransferase /DB_XREF=gi:11863164 /UG=Hs.279865 trans-prenyltransferase /FL=gb:NM_014317.2 gb:AF118395.1	NM_014317		NP_055132
220881_at	0.038535	gb:NM_018606.1 /DEF=Homo sapiens hypothetical protein PRO1787 (PRO1787), mRNA. /FEA=mRNA /GEN=PRO1787 /PROD=hypothetical protein PRO1787 /DB_XREF=gi:8924080 /UG=Hs.283063 hypothetical protein PRO1787 /FL=gb:AF116668.1 gb:NM_018606.1	NM_018606		NP_061076
220924_s_at	1.26E-04	gb:NM_018976.1 /DEF=Homo sapiens amino acid transporter 2 (KIAA1382), mRNA. /FEA=mRNA /GEN=KIAA1382 /PROD=amino acid transporter 2 /DB_XREF=gi:9506836 /UG=Hs.298275 amino acid transporter 2 /FL=gb:NM_018976.1	NM_018976		NP_061849
220926_s_at	3.26E-04	gb:NM_025191.1 /DEF=Homo sapiens hypothetical protein FLJ13033 (FLJ13033), mRNA. /FEA=mRNA /GEN=FLJ13033 /PROD=hypothetical protein FLJ13033 /DB_XREF=gi:13376781 /UG=Hs.301997 hypothetical protein FLJ13033 /FL=gb:NM_025191.1	NM_025191		NP_079467
220939_s_at	0.001116	gb:NM_017743.1 /DEF=Homo sapiens hypothetical protein FLJ20283 (FLJ20283), mRNA. /FEA=mRNA /GEN=FLJ20283 /PROD=hypothetical protein FLJ20283 /DB_XREF=gi:8923261 /UG=Hs.44033 hypothetical protein FLJ20283 /FL=gb:NM_017743.1	NM_017743		NP_569118

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220940_at	2.05E-04	gb:NM_025190.1 /DEF=Homo sapiens KIAA1641 protein (KIAA1641), mRNA. /FEA=mRNA /GEN=KIAA1641 /PROD=hypothetical protein FLJ21281 /DB_XREF=gi:13449272 /UG=Hs.44566 KIAA1641 protein /FL=gb:NM_025190.1	NM_025190		NP_079466
220941_s_at	0.041055	gb:NM_017447.1 /DEF=Homo sapiens hypothetical protein LOC54149 (YG81), mRNA. /FEA=mRNA /GEN=YG81 /PROD=hypothetical protein LOC54149 /DB_XREF=gi:8394546 /UG=Hs.49391 hypothetical protein LOC54149 /FL=gb:NM_017447.1	NM_017447		NP_059143
220945_x_at	1.26E-04	gb:NM_018050.1 /DEF=Homo sapiens hypothetical protein FLJ10298 (FLJ10298), mRNA. /FEA=mRNA /GEN=FLJ10298 /PROD=hypothetical protein FLJ10298 /DB_XREF=gi:8922334 /UG=Hs.5999 hypothetical protein FLJ10298 /FL=gb:NM_018050.1	NM_018050		NP_060520
220947_s_at	7.58E-04	gb:NM_015527.1 /DEF=Homo sapiens DKFZP434P1750 protein (DKFZP434P1750), mRNA. /FEA=mRNA /GEN=DKFZP434P1750 /PROD=DKFZP434P1750 protein /DB_XREF=gi:7661587 /UG=Hs.7274 DKFZP434P1750 protein /FL=gb:NM_015527.1	NM_015527		NP_056342
220949_s_at	1.26E-04	gb:NM_024033.1 /DEF=Homo sapiens hypothetical protein MGC5242 (MGC5242), mRNA. /FEA=mRNA /GEN=MGC5242 /PROD=hypothetical protein MGC5242 /DB_XREF=gi:13162284 /UG=Hs.77365 hypothetical protein MGC5242 /FL=gb:BC000168.2 gb:NM_024033.1	NM_024033		NP_076938

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220951_s_at	0.016495	gb:NM_014576.1 /DEF=Homo sapiens Apobec-1 complementation factor; APOBEC-1 stimulating protein (ACF), mRNA. /FEA=mRNA /GEN=ACF /PROD=Apobec-1 complementation factor; APOBEC-1stimulating protein /DB_XREF=gi:7656852 /UG=Hs.8349 Apobec-1 complementation factor; APOBEC-1 stimulating protein /FL=gb:AF209192.1 gb:NM_014576.1 gb:AF271789.1	NM_014576		NP_620311
220952_s_at	0.041055	gb:NM_019012.1 /DEF=Homo sapiens hypothetical protein (FLJ10667), mRNA. /FEA=mRNA /GEN=FLJ10667 /PROD=hypothetical protein /DB_XREF=gi:9506616 /UG=Hs.86149 phosphoinositol 3-phosphate-binding protein-2 /FL=gb:NM_019012.1	NM_019012		NP_061885
220953_s_at	0.001471	gb:NM_019061.1 /DEF=Homo sapiens hypothetical protein (FLJ20476), mRNA. /FEA=mRNA /GEN=FLJ20476 /PROD=hypothetical protein /DB_XREF=gi:9506678 /UG=Hs.93872 KIAA1682 protein /FL=gb:NM_019061.1	NM_019061		NP_061934
220956_s_at	0.026891	gb:NM_017555.1 /DEF=Homo sapiens hypothetical protein DKFZp434E026 (DKFZp434E026), mRNA. /FEA=mRNA /GEN=DKFZp434E026 /PROD=hypothetical protein DKFZp434E026 /DB_XREF=gi:8922129 /UG=Hs.9659 hypothetical protein DKFZp434E026 /FL=gb:NM_017555.1	NM_017555		NP_542770
220964_s_at	0.010205	gb:NM_030981.1 /DEF=Homo sapiens small GTP-binding protein (RAB1B), mRNA. /FEA=mRNA /GEN=RAB1B /PROD=small GTP-binding protein /DB_XREF=gi:13569961 /FL=gb:NM_030981.1	NM_030981		NP_112243

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220980_s_at	3.26E-04	gb:NM_031284.1 /DEF=Homo sapiens hypothetical protein DKFZp434B195 (DKFZP434B195), mRNA. /FEA=mRNA /GEN=DKFZP434B195 /PROD=hypothetical protein DKFZp434B195 /DB_XREF=gi:13775193 /FL=gb:NM_031284.1	NM_031284		NP_112574
220985_s_at	0.016402	gb:NM_030954.1 /DEF=Homo sapiens hypothetical protein DKFZp564A022 (DKFZP564A022), mRNA. /FEA=mRNA /GEN=DKFZP564A022 /PROD=hypothetical protein DKFZp564A022 /DB_XREF=gi:13569925 /FL=gb:NM_030954.1	NM_030954		NP_112216
220987_s_at	0.016934	gb:NM_030952.1 /DEF=Homo sapiens hypothetical protein DKFZp434J037 (DKFZP434J037), mRNA. /FEA=mRNA /GEN=DKFZP434J037 /PROD=hypothetical protein DKFZp434J037 /DB_XREF=gi:13569921 /FL=gb:NM_030952.1	NM_030952		NP_112214
220995_at	0.00884	gb:NM_014178.1 /DEF=Homo sapiens HSPC156 protein (HSPC156), mRNA. /FEA=mRNA /GEN=HSPC156 /PROD=HSPC156 protein /DB_XREF=gi:7661811 /UG=Hs.99291 HSPC156 protein /FL=gb:AF161505.1 gb:NM_014178.1	NM_014178		NP_054897
220999_s_at	0.016934	gb:NM_030778.1 /DEF=Homo sapiens hypothetical protein PRO1331 (PRO1331), mRNA. /FEA=mRNA /GEN=PRO1331 /PROD=hypothetical protein PRO1331 /DB_XREF=gi:13562115 /FL=gb:NM_030778.1	NM_030778		NP_110405

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221002_s_at	2.05E-04	gb:NM_030927.1 /DEF=Homo sapiens hypothetical protein MGC11352 (MGC11352), mRNA. /FEA=mRNA /GEN=MGC11352 /PROD=hypothetical protein MGC11352 /DB_XREF=gi:13569888 /FL=gb:NM_030927.1	NM_030927		NP_112189
221006_s_at	7.58E-04	gb:NM_030918.1 /DEF=Homo sapiens hypothetical protein My014 (MY014), mRNA. /FEA=mRNA /GEN=MY014 /PROD=hypothetical protein My014 /DB_XREF=gi:13569876 /FL=gb:NM_030918.1	NM_030918		NP_112180
221007_s_at	0.026891	gb:NM_030917.1 /DEF=Homo sapiens hypothetical protein DKFZp586K0717 (DKFZP586K0717), mRNA. /FEA=mRNA /GEN=DKFZP586K0717 /PROD=hypothetical protein DKFZp586K0717 /DB_XREF=gi:13569873 /FL=gb:NM_030917.1	NM_030917		NP_112179
221011_s_at	3.26E-04	gb:NM_030915.1 /DEF=Homo sapiens hypothetical protein DKFZp566J091 (DKFZP566J091), mRNA. /FEA=mRNA /GEN=DKFZP566J091 /PROD=hypothetical protein DKFZp566J091 /DB_XREF=gi:13569871 /FL=gb:NM_030915.1	NM_030915		NP_112177
221012_s_at	0.016934	gb:NM_030912.1 /DEF=Homo sapiens tripartite motif protein TRIM8 (TRIM8), mRNA. /FEA=mRNA /GEN=TRIM8 /PROD=tripartite motif protein TRIM8 /DB_XREF=gi:13569865 /FL=gb:NM_030912.1	NM_030912		NP_112174

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221014_s_at	0.00161	gb:NM_031296.1 /DEF=Homo sapiens hypothetical protein DKFZp434G099 (DKFZP434G099), mRNA. /FEA=mRNA /GEN=DKFZP434G099 /PROD=hypothetical protein DKFZp434G099 /DB_XREF=gi:13786128 /FL=gb:NM_031296.1	NM_031296		NP_112586
221025_x_at	0.041795	gb:NM_031292.1 /DEF=Homo sapiens hypothetical protein DKFZp434G1415 (DKFZP434G1415), mRNA. /FEA=mRNA /GEN=DKFZP434G1415 /PROD=hypothetical protein DKFZp434G1415 /DB_XREF=gi:13775209 /FL=gb:NM_031292.1	NM_031292		NP_112582
221028_s_at	0.011725	gb:NM_030819.1 /DEF=Homo sapiens hypothetical protein MGC11335 (MGC11335), mRNA. /FEA=mRNA /GEN=MGC11335 /PROD=hypothetical protein MGC11335 /DB_XREF=gi:13540615 /FL=gb:NM_030819.1	NM_030819		NP_110446
221036_s_at	1.26E-04	gb:NM_031301.1 /DEF=Homo sapiens hypothetical protein DKFZp564D0372 (DKFZP564D0372), mRNA. /FEA=mRNA /GEN=DKFZP564D0372 /PROD=hypothetical protein DKFZp564D0372 /DB_XREF=gi:13775223 /FL=gb:NM_031301.1	NM_031301		NP_112591
221039_s_at	0.026891	gb:NM_018482.1 /DEF=Homo sapiens KIAA1249 protein (KIAA1249), mRNA. /FEA=mRNA /GEN=KIAA1249 /PROD=uncharacterized gastric protein ZG14P /DB_XREF=gi:8923867 /UG=Hs.10669 KIAA1249 protein /FL=gb:AF222859.1 gb:NM_018482.1	NM_018482		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221041_s_at	0.001109	gb:NM_012434.1 /DEF=Homo sapiens solute carrier family 17 (anionsugar transporter), member 5 (SLC17A5), mRNA. /FEA=mRNA /GEN=SLC17A5 /PROD=solute carrier family 17 (anionsugar transporter), member 5 /DB_XREF=gi:6912665 /UG=Hs.117865 solute carrier family 17 (anionsugar transporter), member 5 /FL=gb:NM_012434.1	NM_012434		NP_036566
221044_s_at	0.005852	gb:NM_021616.1 /DEF=Homo sapiens ring finger protein 21, interferon-responsive (RNF21), mRNA. /FEA=mRNA /GEN=RNF21 /PROD=ring finger protein 21, interferon-responsive /DB_XREF=gi:11055977 /UG=Hs.125300 ring finger protein 21, interferon-responsive /FL=gb:AB039903.1 gb:NM_021616.1	NM_021616		NP_569074
221046_s_at	0.00161	gb:NM_014170.1 /DEF=Homo sapiens HSPC135 protein (HSPC135), mRNA. /FEA=mRNA /GEN=HSPC135 /PROD=HSPC135 protein /DB_XREF=gi:7661795 /UG=Hs.127496 HSPC135 protein /FL=gb:AF161484.1 gb:NM_014170.1	NM_014170		NP_054889
221050_s_at	0.004541	gb:NM_019096.1 /DEF=Homo sapiens GTP binding protein 2 (GTPBP2), mRNA. /FEA=mRNA /GEN=GTPBP2 /PROD=GTP binding protein 2 /DB_XREF=gi:9506762 /UG=Hs.13011 GTP binding protein 2 /FL=gb:NM_019096.1	NM_019096		NP_061969

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221058_s_at	3.26E-04	gb:NM_016326.2 /DEF=Homo sapiens chemokine-like factor 3, alternatively spliced (LOC51192), mRNA. /FEA=mRNA /GEN=LOC51192 /PROD=chemokine-like factor 3, alternatively spliced /DB_XREF=gi:10092611 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM_016326.2 gb:AF135381.2	NM_016326		NP_058647
221059_s_at	7.58E-04	gb:NM_021615.1 /DEF=Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 (CHST6), mRNA. /FEA=mRNA /GEN=CHST6 /PROD=carbohydrate (N-acetylglucosamine 6-O)sulfotransferase 6 /DB_XREF=gi:11055975 /UG=Hs.157439 carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 /FL=gb:AF219990.1 gb:NM_021615.1	NM_021615		NP_067628
221082_s_at	0.003664	gb:NM_022477.1 /DEF=Homo sapiens hypothetical protein FLJ13556 similar to N-myc downstream regulated 3 (FLJ13556), mRNA. /FEA=mRNA /GEN=FLJ13556 /PROD=hypothetical protein FLJ13556 similar to N-mycdownstream regulated 3 /DB_XREF=gi:11968028 /UG=Hs.240615 hypothetical protein FLJ13556 similar to N-myc downstream regulated 3 /FL=gb:NM_022477.1	NM_022477		NP_114402

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221156_x_at	1.26E-04	gb:NM_004748.1 /DEF=Homo sapiens cell cycle progression 8 protein (CPR8), mRNA. /FEA=mRNA /GEN=CPR8 /PROD=cell cycle progression 8 protein /DB_XREF=gi:4758047 /UG=Hs.283753 cell cycle progression 8 protein /FL=gb:AF011794.1 gb:NM_004748.1	NM_004748		NP_004739
221188_s_at	2.05E-04	gb:NM_014430.1 /DEF=Homo sapiens cell death-inducing DFFA-like effector b (CIDEb), mRNA. /FEA=mRNA /GEN=CIDEb /PROD=cell death-inducing DFFA-like effector b /DB_XREF=gi:7656978 /UG=Hs.288835 cell death-inducing DFFA-like effector b /FL=gb:AF190901.1 gb:NM_014430.1	NM_014430		NP_055245
221190_s_at	0.013223	gb:NM_013326.1 /DEF=Homo sapiens colon cancer-associated protein Mic1 (MIC1), mRNA. /FEA=mRNA /GEN=MIC1 /PROD=colon cancer-associated protein Mic1 /DB_XREF=gi:7019454 /UG=Hs.289080 colon cancer-associated protein Mic1 /FL=gb:AF143536.1 gb:NM_013326.1	NM_013326		NP_037458
221194_s_at	0.016495	gb:NM_016125.1 /DEF=Homo sapiens PTD016 protein (LOC51136), mRNA. /FEA=mRNA /GEN=LOC51136 /PROD=PTD016 protein /DB_XREF=gi:7705842 /UG=Hs.30154 PTD016 protein /FL=gb:AF100745.1 gb:NM_016125.1	NM_016125		NP_057209
221196_x_at	0.00945	gb:NM_024332.1 /DEF=Homo sapiens c6.1A (C6.1A), mRNA. /FEA=mRNA /GEN=C6.1A /PROD=c6.1A /DB_XREF=gi:13236582 /UG=Hs.301927 c6.1A /FL=gb:BC002999.1 gb:NM_024332.1	NM_024332		NP_077308

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221210_s_at	0.007779	gb:NM_030769.1 /DEF=Homo sapiens hypothetical protein similar to swine acylneuraminate lyase (C1ORF13), mRNA. /FEA=mRNA /GEN=C1ORF13 /PROD=hypothetical protein similar to swineacylneuraminate lyase /DB_XREF=gi:13540532 /FL=gb:NM_030769.1	NM_030769		NP_110396
221213_s_at	0.026884	gb:NM_017661.1 /DEF=Homo sapiens hypothetical protein FLJ20086 (FLJ20086), mRNA. /FEA=mRNA /GEN=FLJ20086 /PROD=hypothetical protein FLJ20086 /DB_XREF=gi:8923096 /UG=Hs.46821 hypothetical protein FLJ20086 /FL=gb:NM_017661.1	NM_017661		NP_060131
221214_s_at	5.03E-04	gb:NM_015537.1 /DEF=Homo sapiens DKFZP586J1624 protein (DKFZP586J1624), mRNA. /FEA=mRNA /GEN=DKFZP586J1624 /PROD=DKFZP586J1624 protein /DB_XREF=gi:7661679 /UG=Hs.4964 DKFZP586J1624 protein /FL=gb:NM_015537.1	NM_015537		NP_056352
221218_s_at	0.00228	gb:NM_022445.1 /DEF=Homo sapiens mouse thiamin pyrophosphokinase homolog (TPK1), mRNA. /FEA=mRNA /GEN=TPK1 /PROD=mouse thiamin pyrophosphokinase homolog /DB_XREF=gi:11990617 /UG=Hs.58715 thiamine pyrophosphokinase /FL=gb:NM_022445.1	NM_022445		NP_071890
221234_s_at	0.00161	gb:NM_021813.1 /DEF=Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription factor 2 (BACH2), mRNA. /FEA=mRNA /GEN=BACH2 /PROD=BTB and CNC homology 1, basic leucine zippertranscription factor 2 /DB_XREF=gi:13540489 /FL=gb:NM_021813.1	NM_021813		NP_068585

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221235_s_at	0.001116	gb:NM_030825.1 /DEF=Homo sapiens hypothetical protein FLJ22294 (FLJ22294), mRNA. /FEA=mRNA /GEN=FLJ22294 /PROD=hypothetical protein FLJ22294 /DB_XREF=gi:13540497 /FL=gb:NM_030825.1	NM_030825		
221249_s_at	0.003171	gb:NM_030802.1 /DEF=Homo sapiens CEBP-induced protein (LOC81558), mRNA. /FEA=mRNA /GEN=LOC81558 /PROD=CEBP-induced protein /DB_XREF=gi:13540589 /FL=gb:NM_030802.1	NM_030802		NP_110429
221257_x_at	3.26E-04	gb:NM_030793.1 /DEF=Homo sapiens hypothetical protein SP329 (SP329), mRNA. /FEA=mRNA /GEN=SP329 /PROD=hypothetical protein SP329 /DB_XREF=gi:13540573 /FL=gb:NM_030793.1	NM_030793		NP_110420
221263_s_at	7.58E-04	gb:NM_031287.1 /DEF=Homo sapiens hypothetical protein MGC3133 (MGC3133), mRNA. /FEA=mRNA /GEN=MGC3133 /PROD=hypothetical protein MGC3133 /DB_XREF=gi:13775199 /FL=gb:NM_031287.1	NM_031287		NP_112577
221264_s_at	1.26E-04	gb:NM_031214.1 /DEF=Homo sapiens hypothetical protein (AF311304), mRNA. /FEA=mRNA /GEN=AF311304 /PROD=hypothetical protein /DB_XREF=gi:13654285 /FL=gb:NM_031214.1	NM_031214		NP_112491
221269_s_at	0.041055	gb:NM_031286.1 /DEF=Homo sapiens SH3BGRL3-like protein (SH3BGRL3), mRNA. /FEA=mRNA /GEN=SH3BGRL3 /PROD=SH3BGRL3-like protein /DB_XREF=gi:13775197 /FL=gb:NM_031286.1	NM_031286		NP_112576

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221332_at	0.041795	gb:NM_005448.1 /DEF=Homo sapiens bone morphogenetic protein 15 (BMP15), mRNA. /FEA=CDS /GEN=BMP15 /PROD=bone morphogenetic protein 15 precursor /DB_XREF=gi:4885096 /UG=Hs.247820 bone morphogenetic protein 15 /FL=gb:NM_005448.1	NM_005448		NP_005439
221381_s_at	1.26E-04	gb:NM_006792.2 /DEF=Homo sapiens mortality factor 4 (MORF4), mRNA. /FEA=CDS /GEN=MORF4 /PROD=mortality factor 4 /DB_XREF=gi:6996927 /UG=Hs.251396 mortality factor 4 /FL=gb:NM_006792.2	NM_006792		
221419_s_at	0.00228	gb:NM_013307.1 /DEF=Homo sapiens non-functional folate binding protein (HSAF000381), mRNA. /FEA=CDS /GEN=HSAF000381 /PROD=non-functional folate binding protein /DB_XREF=gi:7019412 /FL=gb:NM_013307.1	NM_013307		
221421_s_at	0.015246	gb:NM_030955.1 /DEF=Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 12 (ADAMTS12), mRNA. /FEA=CDS /GEN=ADAMTS12 /PROD=a disintegrin-like and metalloprotease(reprolysin type) with thrombospondin type 1 motif, 12 /DB_XREF=gi:13569927 /FL=gb:NM_030955.1	NM_030955		NP_112217
221425_s_at	2.76E-04	gb:NM_030940.1 /DEF=Homo sapiens hypothetical protein MGC4276 similar to CG8198 (MGC4276), mRNA. /FEA=CDS /GEN=MGC4276 /PROD=hypothetical protein MGC4276 similar to CG8198 /DB_XREF=gi:13569910 /FL=gb:NM_030940.1	NM_030940		NP_112202

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221428_s_at	0.007779	gb:NM_030921.1 /DEF=Homo sapiens hypothetical protein DC42 (DC42), mRNA. /FEA=CDS /GEN=DC42 /PROD=hypothetical protein DC42 /DB_XREF=gi:13569880 /FL=gb:NM_030921.1	NM_030921		NP_112183
221430_s_at	0.007779	gb:NM_030963.1 /DEF=Homo sapiens hypothetical protein DKFZp434O1427 (DKFZP434O1427), mRNA. /FEA=CDS /GEN=DKFZP434O1427 /PROD=hypothetical protein DKFZp434O1427 /DB_XREF=gi:13624336 /FL=gb:NM_030963.1	NM_030963		NP_112225
221437_s_at	0.025464	gb:NM_031280.1 /DEF=Homo sapiens mitochondrial ribosomal protein S15 (MRPS15), mRNA. /FEA=CDS /GEN=MRPS15 /PROD=mitochondrial ribosomal protein S15 /DB_XREF=gi:13775191 /FL=gb:NM_031280.1	NM_031280		NP_112570
221449_s_at	5.07E-04	gb:NM_030790.1 /DEF=Homo sapiens hypothetical protein CDA08 (CDA08), mRNA. /FEA=CDS /GEN=CDA08 /PROD=hypothetical protein CDA08 /DB_XREF=gi:13540566 /FL=gb:NM_030790.1	NM_030790		NP_110417
221452_s_at	3.26E-04	gb:NM_030969.1 /DEF=Homo sapiens hypothetical protein MGC1223 (MGC1223), mRNA. /FEA=CDS /GEN=MGC1223 /PROD=hypothetical protein MGC1223 /DB_XREF=gi:13624338 /FL=gb:NM_030969.1	NM_030969		NP_112231
221471_at	1.26E-04	tumor differentially expressed 1	AW173623	Hs.272168	NP_006802
221472_at	1.26E-04	tumor differentially expressed 1	AW173623	Hs.272168	NP_006802

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221473_x_at	0.003171	gb:U49188.1 /DEF=Human placenta (Diff33) mRNA, complete cds. /FEA=mRNA /GEN=Diff33 /DB_XREF=gi:1293562 /UG=Hs.272168 tumor differentially expressed 1 /FL=gb:U49188.1 gb:AF112227.1 gb:AF153979.1 gb:NM_006811.1	U49188		NP_006802
221474_at	0.013223	gb:U26162.1 /DEF=Human myosin regulatory light chain mRNA, complete cds. /FEA=mRNA /PROD=myosin regulatory light chain /DB_XREF=gi:829622 /UG=Hs.180224 death-associated protein 6 /FL=gb:BC004994.1 gb:D50372.1 gb:U26162.1	U26162		NP_291024
221475_s_at	0.026891	Consensus includes gb:NM_002948.1 /DEF=Homo sapiens ribosomal protein L15 (RPL15), mRNA. /FEA=CDS /GEN=RPL15 /PROD=ribosomal protein L15 /DB_XREF=gi:4506602 /UG=Hs.74267 ribosomal protein L15 /FL=gb:AF279903.1 gb:L25899.1 gb:NM_002948.1	NM_002948		NP_002939
221477_s_at	1.26E-04	superoxide dismutase 2, mitochondrial	BF575213	Hs.372783	
221478_at	2.05E-04	Consensus includes gb:AL132665.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566E034 (from clone DKFZp566E034); complete cds. /FEA=mRNA /DB_XREF=gi:6137021 /UG=Hs.132955 BCL2adenovirus E1B 19kD-interacting protein 3-like /FL=gb:AF060922.1 gb:AB004788.1 gb:AF067396.1 gb:NM_004331.1 gb:AL132665.1	AL132665		NP_004322

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221479_s_at	2.05E-04	gb:AF060922.1 /DEF=Homo sapiens clone 016a05 My020 protein mRNA, complete cds. /FEA=mRNA /PROD=My020 protein /DB_XREF=gi:12001981 /UG=Hs.132955 BCL2adenovirus E1B 19kD-interacting protein 3-like /FL=gb:AF060922.1 gb:AB004788.1 gb:AF067396.1 gb:NM_004331.1 gb:AL132665.1	AF060922		NP_004322
221480_at	0.019879	heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kDa)	BG180941	Hs.303627	NP_112738
221482_s_at	0.001116	gb:BC003418.1 /DEF=Homo sapiens, cyclic AMP phosphoprotein, 19 kD, clone MGC:5468, mRNA, complete cds. /FEA=mRNA /PROD=cyclic AMP phosphoprotein, 19 kD /DB_XREF=gi:13097335 /UG=Hs.7351 cyclic AMP phosphoprotein, 19 kD /FL=gb:BC003418.1 gb:AF084555.1	BC003418		NP_006619
221484_at	0.007779	Consensus includes gb:BF691447 /FEA=EST /DB_XREF=gi:11976855 /DB_XREF=est:602247615F1 /CLONE=IMAGE:4332866 /UG=Hs.107526 UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5 /FL=gb:AB004550.1 gb:AF038663.1 gb:NM_004776.1	NM_004776		NP_004767

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221485_at	3.26E-04	Consensus includes gb:AL035683 /DEF=Human DNA sequence from clone RP5-1063B2 on chromosome 20q13.1-13.2. Contains the 3 part of the gene for Beta-1,4-galactosyltransferase, ESTs, STSs and GSSs /FEA=mRNA /DB_XREF=gi:7288039 /UG=Hs.107526 UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5 /FL=gb:AB004550.1 gb:AF038663.1 gb:NM_004776.1	NM_004776		NP_004767
221486_at	0.021244	gb:AF067170.1 /DEF=Homo sapiens alpha endosulfine mRNA, complete cds. /FEA=mRNA /PROD=alpha endosulfine /DB_XREF=gi:4894373 /UG=Hs.111680 endosulfine alpha /FL=gb:AF067170.1 gb:AF157510.1	AF067170		NP_004427
221489_s_at	0.021312	Consensus includes gb:W48843 /FEA=EST /DB_XREF=gi:1337022 /DB_XREF=est:zc44a09.s1 /CLONE=IMAGE:325144 /UG=Hs.285814 sprouty (Drosophila) homolog 4 /FL=gb:AF227517.1	AF227517		
221490_at	1.41E-04	gb:AL136733.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N1010 (from clone DKFZp434N1010); complete cds. /FEA=mRNA /GEN=DKFZp434N1010 /PROD=hypothetical protein /DB_XREF=gi:12052984 /UG=Hs.75425 ubiquitin associated protein /FL=gb:AL136733.1 gb:AF222043.2 gb:NM_016525.2	AL136733		NP_057609

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221492_s_at	1.26E-04	gb:AF202092.1 /DEF=Homo sapiens PC3-96 mRNA, complete cds. /FEA=mRNA /PROD=PC3-96 /DB_XREF=gi:11493699 /UG=Hs.26367 PC3-96 protein /FL=gb:AF202092.1	AF202092		NP_071933
221497_x_at	0.00161	gb:BC005369.1 /DEF=Homo sapiens, chromosome 1 open reading frame 12, clone MGC:12484, mRNA, complete cds. /FEA=mRNA /PROD=chromosome 1 open reading frame 12 /DB_XREF=gi:13529208 /UG=Hs.6523 chromosome 1 open reading frame 12 /FL=gb:AF229245.1 gb:AF277176.1 gb:NM_022051.1 gb:BC005369.1	BC005369		NP_071334
221498_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	BF939727	Hs.409278	NP_112180
221500_s_at	0.013223	Consensus includes gb:BE782754 /FEA=EST /DB_XREF=gi:10203952 /DB_XREF=est:601471903F1 /CLONE=IMAGE:3874921 /UG=Hs.102178 syntaxin 16 /FL=gb:AF008936.1	AF008936		NP_003754
221504_s_at	1.41E-04	gb:AF112204.1 /DEF=Homo sapiens Vacuolar proton pump subunit SFD alpha isoform mRNA, complete cds. /FEA=mRNA /PROD=Vacuolar proton pump subunit SFD alpha isoform /DB_XREF=gi:6563195 /UG=Hs.19575 CGI-11 protein /FL=gb:AF298777.1 gb:AF132945.1 gb:AF112204.1 gb:NM_015941.1	AF112204		NP_057025
221505_at	1.26E-04	leucine-rich acidic nuclear protein like	AW612574	Hs.71331	NP_112182
221506_s_at	0.020412	karyopherin beta 2b, transportin	BG258639	Hs.278378	
221507_at	6.68E-04	karyopherin beta 2b, transportin	BG258639	Hs.278378	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221511_x_at	1.26E-04	Consensus includes gb:AB033080.1 /DEF=Homo sapiens mRNA for KIAA1254 protein, partial cds. /FEA=mRNA /GEN=KIAA1254 /PROD=KIAA1254 protein /DB_XREF=gi:6330892 /UG=Hs.82506 KIAA1254 protein /FL=gb:AF212228.1	AB033080		NP_004739
221515_s_at	0.006295	gb:BC001214.1 /DEF=Homo sapiens, CGI-68 protein, clone MGC:682, mRNA, complete cds. /FEA=mRNA /PROD=CGI-68 protein /DB_XREF=gi:12654744 /UG=Hs.8054 CGI-68 protein /FL=gb:BC001214.1 gb:AF151826.1 gb:NM_016015.1	BC001214		NP_057393
221516_s_at	0.001116	gb:BC002587.1 /DEF=Homo sapiens, hypothetical protein, clone MGC:1067, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:12803520 /UG=Hs.83869 hypothetical protein /FL=gb:BC002587.1	BC002587		NP_061881
221522_at	0.041055	gb:AL136784.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L0718 (from clone DKFZp434L0718); complete cds. /FEA=mRNA /GEN=DKFZp434L0718 /PROD=hypothetical protein /DB_XREF=gi:12053080 /UG=Hs.59236 Homo sapiens mRNA; cDNA DKFZp434L0718 (from clone DKFZp434L0718); complete cds /FL=gb:AL136784.1	AL136784		NP_115515

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221523_s_at	1.26E-04	Consensus includes gb:AL138717 /DEF=Human DNA sequence from clone RP11-11D8 on chromosome 6 Contains the 5 end of the gene for a yeast ubiquitin conjugating enzyme UBC6 homolog, the gene for a possible GTP binding protein, a NACA (nascent-polypeptide-associated complex alpha polype... /FEA=mRNA_2 /DB_XREF=gi:8894207 /UG=Hs.238679 Rag D protein /FL=gb:NM_021244.1 gb:AF272036.1 gb:BC003088.1	AL138717		
221524_s_at	4.42E-04	gb:AF272036.1 /DEF=Homo sapiens Rag D mRNA, complete cds. /FEA=mRNA /PROD=Rag D /DB_XREF=gi:11181619 /UG=Hs.238679 Rag D protein /FL=gb:NM_021244.1 gb:AF272036.1 gb:BC003088.1	AF272036		NP_067067
221528_s_at	6.42E-04	gb:BC000143.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ11656, clone MGC:5247, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ11656 /DB_XREF=gi:12652782 /UG=Hs.96560 hypothetical protein FLJ11656 /FL=gb:BC000143.1	BC000143		NP_573403

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221539_at	7.58E-04	gb:AB044548.1 /DEF=Homo sapiens POKcl.6 mRNA for eukaryotic translation initiation factor 4E binding protein 1, complete cds. /FEA=mRNA /GEN=POKcl.6° /PROD=eukaryotic translation initiation factor 4Ebinding protein 1 /DB_XREF=gi:11527778 /UG=Hs.71819 eukaryotic translation initiation factor 4E binding protein 1 /FL=gb:AB044548.1 gb:BC004459.1 gb:NM_004095.1 gb:L36055.1	AB044548		NP_004086
221542_s_at	0.027418	SRY (sex determining region Y)-box 5	T90773	Hs.115232	NP_009106
221543_s_at	1.26E-04	gb:AL442077.1 /DEF=Homo sapiens mRNA; cDNA DKFZp667H242 (from clone DKFZp667H242); complete cds. /FEA=mRNA /GEN=DKFZp667H242 /PROD=hypothetical protein /DB_XREF=gi:10241715 /UG=Hs.125849 chromosome 8 open reading frame 2 /FL=gb:AL442077.1 gb:NM_007175.1	AL442077		NP_009106
221547_at	1.26E-04	gb:BC000794.1 /DEF=Homo sapiens, pre-mRNA splicing factor similar to S. cerevisiae Prp18, clone MGC:5075, mRNA, complete cds. /FEA=mRNA /PROD=pre-mRNA splicing factor similar to S.cerevisiae Prp18 /DB_XREF=gi:12653992 /UG=Hs.155244 pre-mRNA processing factor 18 /FL=gb:BC000794.1 gb:U51990.1 gb:NM_003675.1	BC000794		NP_003666

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221550_at	0.048473	gb:BC002382.1 /DEF=Homo sapiens, COX15 (yeast) homolog, cytochrome c oxidase assembly protein, clone MGC:8634, mRNA, complete cds. /FEA=mRNA /PROD=COX15 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:12803156 /UG=Hs.226581 COX15 (yeast) homolog, cytochrome c oxidase assembly protein /FL=gb:BC002382.1 gb:AF026850.1	BC002382		NP_510870
221553_at	0.013223	gb:AL136636.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564K142 (from clone DKFZp564K142); complete cds. /FEA=mRNA /GEN=DKFZp564K142 /PROD=hypothetical protein /DB_XREF=gi:12052797 /UG=Hs.323562 Homo sapiens mRNA; cDNA DKFZp564K142 (from clone DKFZp564K142); complete cds /FL=gb:AL136636.1	AL136636		NP_115497
221558_s_at	0.016934	gb:AF288571.1 /DEF=Homo sapiens lymphoid enhancer factor-1 (LEF1) mRNA, complete cds. /FEA=mRNA /GEN=LEF1 /PROD=lymphoid enhancer factor-1 /DB_XREF=gi:9858157 /UG=Hs.44865 lymphoid enhancer binding factor-1 /FL=gb:AF198532.1 gb:NM_016269.1 gb:AF288571.1	AF288571		NP_057353

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221561_at	0.033381	gb:L21934.2 /DEF=Homo sapiens acyl-coenzyme A: cholesterol acyltransferase mRNA, complete cds. /FEA=mRNA /PROD=acyl-coenzyme A: cholesterol acyltransferase /DB_XREF=gi:4878021 /UG=Hs.14553 sterol O-acyltransferase (acyl-Coenzyme A: cholesterol acyltransferase) 1 /FL=gb:NM_003101.1 gb:L21934.2	L21934		NP_003092
221564_at	0.010205	HMT1 hnRNP methyltransferase-like 1 (<i>S. cerevisiae</i>)	AL570294	Hs.235887	NP_001526
221571_at	5.03E-04	TNF receptor-associated factor 3	AI721219	Hs.297660	NP_663778
221581_s_at	0.007779	gb:AF257135.1 /DEF=Homo sapiens WBSCR15 protein (WBSCR15) mRNA, complete cds. /FEA=mRNA /GEN=WBSCR15 /PROD=WBSCR15 protein /DB_XREF=gi:9651998 /UG=Hs.56607 Williams-Beuren syndrome chromosome region 5 /FL=gb:NM_022040.1 gb:BC001609.1 gb:AF257135.1	AF257135		NP_115853
221588_x_at	0.010205	Consensus includes gb:AI640855 /FEA=EST /DB_XREF=gi:4703964 /DB_XREF=est:wa27c11.x1 /CLONE=IMAGE:2299316 /UG=Hs.293970 methylmalonate-semialdehyde dehydrogenase /FL=gb:AF130089.1	AF130089		
221596_s_at	0.02145	gb:AL136619.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564O0523 (from clone DKFZp564O0523); complete cds. /FEA=mRNA /GEN=DKFZp564O0523 /PROD=hypothetical protein /DB_XREF=gi:12052763 /UG=Hs.21590 Homo sapiens mRNA; cDNA DKFZp564O0523 (from clone DKFZp564O0523); complete cds /FL=gb:AL136619.1	AL136619		NP_115496

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221597_s_at	0.033381	gb:BC003080.1 /DEF=Homo sapiens, Similar to HSPC171 protein, clone MGC:770, mRNA, complete cds. /FEA=mRNA /PROD=Similar to HSPC171 protein /DB_XREF=gi:13111781 /UG=Hs.279593 HSPC171 protein /FL=gb:BC003080.1	BC003080		NP_054906
221601_s_at	0.001618	regulator of Fas-induced apoptosis	AI084226	Hs.58831	NP_005440
221613_s_at	3.26E-04	gb:AL136598.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F2116 (from clone DKFZp564F2116); complete cds. /FEA=mRNA /GEN=DKFZp564F2116 /PROD=hypothetical protein /DB_XREF=gi:13276696 /UG=Hs.83954 protein associated with PRK1 /FL=gb:AL136598.1	AL136598		NP_061879
221616_s_at	0.041055	Consensus includes gb:AF077053.1 /DEF=Homo sapiens neuronal cell death-related protein mRNA, complete cds. /FEA=mRNA /PROD=neuronal cell death-related protein /DB_XREF=gi:4689153 /UG=Hs.171723 neuronal cell death-related protein /FL=gb:AF077053.1 gb:NM_015975.1 gb:AF220509.1	AF077053		NP_057059
221620_s_at	3.26E-04	gb:AF061264.1 /DEF=Homo sapiens brain my025 mRNA, complete cds. /FEA=mRNA /PROD=brain my025 /DB_XREF=gi:12001991 /UG=Hs.40065 hypothetical protein MGC4825 /FL=gb:AF061264.1	AF061264		NP_077027

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221637_s_at	0.003171	gb:BC001434.1 /DEF=Homo sapiens, clone MGC:2477, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:2477) /DB_XREF=gi:12655158 /UG=Hs.9061 hypothetical protein MGC2477 /FL=gb:BC001434.1	BC001434		NP_077004
221647_s_at	0.02145	gb:AL136935.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586O0222 (from clone DKFZp586O0222); complete cds. /FEA=mRNA /GEN=DKFZp586O0222 /PROD=hypothetical protein /DB_XREF=gi:12053364 /UG=Hs.7393 hypothetical protein from EUROIMAGE 1987170 /FL=gb:AL136935.1	AL136935		NP_068751
221654_s_at	5.03E-04	gb:AF077040.1 /DEF=Homo sapiens SIH003 mRNA, complete cds. /FEA=mRNA /PROD=SIH003 /DB_XREF=gi:4689127 /UG=Hs.251636 ubiquitin specific protease 3 /FL=gb:AF077040.1	AF077040		NP_006528
221658_s_at	0.023856	gb:AF269133.1 /DEF=Homo sapiens novel interleukin receptor (NILR) mRNA, complete cds. /FEA=mRNA /GEN=NILR /PROD=novel interleukin receptor /DB_XREF=gi:10801190 /UG=Hs.210546 interleukin 21 receptor /FL=gb:AF269133.1	AF269133		NP_851565
221664_s_at	0.029453	gb:AF154005.1 /DEF=Homo sapiens junction adhesion molecule mRNA, complete cds. /FEA=mRNA /PROD=junction adhesion molecule /DB_XREF=gi:5457118 /UG=Hs.286218 junctional adhesion molecule /FL=gb:AF154005.1	AF154005		NP_653087

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221666_s_at	7.58E-04	gb:BC004470.1 /DEF=Homo sapiens, clone MGC:10332, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10332) /DB_XREF=gi:13325315 /UG=Hs.71869 apoptosis-associated speck-like protein containing a CARD /FL=gb:BC004470.1	BC004470		NP_660184
221669_s_at	0.026891	gb:BC001964.1 /DEF=Homo sapiens, acyl-Coenzyme A dehydrogenase family, member 8, clone MGC:4966, mRNA, complete cds. /FEA=mRNA /PROD=acyl-Coenzyme A dehydrogenase family, member 8 /DB_XREF=gi:12805020 /UG=Hs.14791 acyl-Coenzyme A dehydrogenase family, member 8 /FL=gb:BC001964.1	BC001964		NP_055199
221670_s_at	0.006966	gb:AF156888.1 /DEF=Homo sapiens LIM homeobox protein 3 isoform a (LHX3) mRNA, complete cds. /FEA=mRNA /GEN=LHX3 /PROD=LIM homeobox protein 3 isoform a /DB_XREF=gi:7110142 /UG=Hs.148427 LIM homeobox protein 3 /FL=gb:AF156888.1	AF156888		NP_835258
221675_s_at	1.26E-04	gb:AF195624.1 /DEF=Homo sapiens cholinephosphotransferase 1 beta mRNA, complete cds. /FEA=mRNA /PROD=cholinephosphotransferase 1 beta /DB_XREF=gi:9502012 /UG=Hs.171889 cholinephosphotransferase 1 /FL=gb:AF195624.1	AF195624		NP_064629
221691_x_at	0.005852	gb:AB042278.1 /DEF=Homo sapiens mRNA for nucleophosminB23.2, complete cds. /FEA=mRNA /PROD=nucleophosminB23.2 /DB_XREF=gi:13536990 /FL=gb:AB042278.1	AB042278		NP_002511

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221702_s_at	0.041055	gb:AF353992.1 /DEF=Homo sapiens BBP-like protein 2 (BLP2) mRNA, complete cds. /FEA=mRNA /GEN=BLP2 /PROD=BBP-like protein 2 /DB_XREF=gi:13625462 /FL=gb:AF353992.1	AF353992		NP_510883
221705_s_at	0.018973	gb:BC005934.1 /DEF=Homo sapiens, hypothetical protein FLJ21168, clone MGC:14549, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein FLJ21168 /DB_XREF=gi:13543553 /FL=gb:BC005934.1	BC005934		NP_079349
221717_at	0.003638	gb:L25664.1 /DEF=Human interferon mRNA, complete cds. /FEA=CDS /PROD=interferon /DB_XREF=gi:479010 /UG=Hs.247726 Human interferon mRNA, complete cds /FL=gb:L25664.1	L25664		
221724_s_at	3.26E-04	gb:AF200738.1 /DEF=Homo sapiens C-type lectin DDB27 short form (DDB27) mRNA, complete cds. /FEA=CDS /GEN=DDB27 /PROD=C-type lectin DDB27 short form /DB_XREF=gi:11493653 /UG=Hs.115515 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /FL=gb:AF200738.1	AF200738		NP_057268
221725_at	3.26E-04	Consensus includes gb:A1962978 /FEA=EST /DB_XREF=gi:5755691 /DB_XREF=est:wt25f05.x1 /CLONE=IMAGE:2508513 /UG=Hs.288908 WAS protein family, member 2 /FL=gb:NM_006990.1 gb:AB026542.1	NM_006990		NP_008921

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221731_x_at	0.003171	Consensus includes gb:BF218922 /FEA=EST /DB_XREF=gi:11112418 /DB_XREF=est:601885091F1 /CLONE=IMAGE:4103447 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican)	J02814		NP_004376
221732_at	0.029713	Consensus includes gb:AK026161.1 /DEF=Homo sapiens cDNA: FLJ22508 fis, clone HRC11772. /FEA=mRNA /DB_XREF=gi:10438922 /UG=Hs.8859 Homo sapiens, Similar to RIKEN cDNA 5830420C20 gene, clone IMAGE:3633379, mRNA, partial cds	AK026161		NP_620148
221735_at	0.046687	WD repeat endosomal protein	H04342	Hs.109778	NP_065890
221736_at	2.05E-04	KIAA1219 protein	BG236163	Hs.25431	
221738_at	0.041055	KIAA1219 protein	BG236163	Hs.25431	
221740_x_at	0.04734	Homo sapiens cDNA FLJ34306 fis, clone FEBRA2008086	BG249885	Hs.339808	NP_055649
221741_s_at	7.58E-04	match: proteins: Tr:Q9Y5A9 Tr:Q9VBZ5 Tr:O77335 Tr:Q9Y172 Tr:O04503; Human DNA sequence from clone RP5-963E22 on chromosome 20 Contains the 3' end of a novel gene similar to NY-REN-2 Antigen, 5 CpG islands, ESTs, STSs and GSSs, complete sequence.	AL096828		
221748_s_at	7.22E-04	tensin	AL046979	Hs.351432	
221749_at	7.58E-04	hypothetical protein FLJ31657	AU157915	Hs.409076	NP_689971
221752_at	1.26E-04	Consensus includes gb:AL041728 /FEA=EST /DB_XREF=gi:5421076 /DB_XREF=est:DKFZp434O0717_s1 /CLONE=DKFZp434O0717 /UG=Hs.60377 KIAA1298 protein	AB037719		NP_061857
221757_at	0.002671	Consensus includes gb:BE042976 /FEA=EST /DB_XREF=gi:8360029 /DB_XREF=est:ho31d03.x1 /CLONE=IMAGE:3038981 /UG=Hs.26670 Human PAC clone RP3-515N1 from 22q11.2-q22	AC002073		
221761_at	1.26E-04	adenylosuccinate synthase	AA628948	Hs.90011	NP_001117

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221765_at	0.001747	UDP-glucose ceramide glucosyltransferase	BF970427	Hs.395863	NP_003349
221766_s_at	0.016934	Consensus includes gb:AW246673 /FEA=EST /DB_XREF=gi:6589666 /DB_XREF=est:2821951.3prime /CLONE=IMAGE:2821951 /UG=Hs.10784 hypothetical protein FLJ20037	AL078599		
221767_x_at	2.05E-04	Consensus includes gb:AA515560 /FEA=EST /DB_XREF=gi:2255160 /DB_XREF=est:ni42c09.s1 /CLONE=IMAGE:979504 /UG=Hs.283473 hypothetical protein PRO2900	AK026138		
221774_x_at	0.041055	transcription factor (p38 interacting protein)	AW003334	Hs.396316	NP_060039
221776_s_at	0.010205	bromodomain containing 7	AI885109	Hs.279762	NP_037395
221778_at	7.58E-04	KIAA1718 protein	BE217882	Hs.222707	
221786_at	1.26E-04	Consensus includes gb:BF197222 /FEA=EST /DB_XREF=gi:11085906 /DB_XREF=est:7m88b07.x1 /CLONE=IMAGE:3561949 /UG=Hs.12342 Homo sapiens clone 24538 mRNA sequence	AF055030		NP_579866
221787_at	7.58E-04	Consensus includes gb:BF431618 /FEA=EST /DB_XREF=gi:11443732 /DB_XREF=est:7o16e03.x1 /CLONE=IMAGE:3574349 /UG=Hs.12342 Homo sapiens clone 24538 mRNA sequence	AF055030		NP_579866
221790_s_at	0.009689	LDL receptor adaptor protein	AL545035	Hs.184482	NP_056442
221798_x_at	0.003171	cadherin 1, type 1, E-cadherin (epithelial)	AI183766	Hs.356360	NP_002943
221802_s_at	0.005339	KIAA1598 protein	AU157109	Hs.23740	NP_060800
221803_s_at	1.26E-04	nuclear receptor binding factor-2	AA883074	Hs.27181	NP_110386
221819_at	0.007779	RAB35, member RAS oncogene family	BF791960	Hs.94308	NP_006852

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221820_s_at	5.03E-04	Consensus includes gb:AK024102.1 /DEF=Homo sapiens cDNA FLJ14040 fis, clone HEMBA1005513, weakly similar to MALES-ABSENT ON THE FIRST PROTEIN (EC 2.3.1.). /FEA=mRNA /DB_XREF=gi:10436399 /UG=Hs.42343 member of MYST family histone acetyl transferases, homolog of Drosophila MOF	AK024102		NP_115564
221823_at	5.64E-04	hypothetical gene supported by AF038182; BC009203	AL565741	Hs.25925	NP_149988
221824_s_at	0.001116	hypothetical protein MGC26766	AA770170	Hs.395949	NP_659458
221829_s_at	0.001116	karyopherin (importin) beta 2	AI307759	Hs.168075	NP_694858
221830_at	1.26E-04	ESTs	AI302106	Hs.409145	NP_066361
221832_s_at	0.043705	Consensus includes gb:AV741657 /FEA=EST /DB_XREF=gi:10859238 /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1	AK025345		
221840_at	0.00228	ESTs	AA775177	Hs.408754	NP_569119
221853_s_at	0.004341	pM5 protein	N39536	Hs.227823	NP_775885
221856_s_at	0.003171	Consensus includes gb:AL561943 /FEA=EST /DB_XREF=gi:12909874 /DB_XREF=est:AL561943 /CLONE=CS0DB002YO04 (3 prime) /UG=Hs.3346 hypothetical protein FLJ11280	AB037811		
221864_at	0.02145	hypothetical protein MGC13024	AW517464	Hs.333488	NP_689501
221867_at	0.001911	hypothetical protein FLJ31821	BF436315	Hs.351575	NP_694574
221875_x_at	0.007779	major histocompatibility complex, class I, F	AW514210	Hs.377850	NP_061823
221892_at	0.009682	Consensus includes gb:AK024548.1 /DEF=Homo sapiens cDNA: FLJ20895 fis, clone ADKA03483. /FEA=mRNA /DB_XREF=gi:10436853 /UG=Hs.285737 Homo sapiens cDNA: FLJ20895 fis, clone ADKA03483	AK024548		
221896_s_at	0.003171	likely ortholog of mouse hypoxia induced gene 1	BE739519	Hs.7917	NP_054775
221919_at	0.033381	heterogeneous nuclear ribonucleoprotein A1	AW450929	Hs.249495	NP_112420
221932_s_at	0.005379	clone FLB4739	AA133341	Hs.294083	NP_057501

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221943_x_at	0.00161	ribosomal protein L38	AW303136	Hs.2017	NP_000990
221952_x_at	5.03E-04	Consensus includes gb:AB037814.1 /DEF=Homo sapiens mRNA for KIAA1393 protein, partial cds. /FEA=mRNA /GEN=KIAA1393 /PROD=KIAA1393 protein /DB_XREF=gi:7243166 /UG=Hs.33317 KIAA1393 protein	AB037814		
221954_at	0.013223	chromosome 20 open reading frame 111	AA160474	Hs.75798	NP_057554
221960_s_at	0.002354	RAB2, member RAS oncogene family	AI189609	Hs.78305	NP_002856
221962_s_at	0.03002	ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)	AI829920	Hs.409297	NP_003335
221978_at	0.001116	major histocompatibility complex, class I, F	BE138825	Hs.377850	NP_061823
221983_at	1.26E-04	hypothetical protein MGC3035	AL040896	Hs.22412	NP_077269
221984_s_at	1.26E-04	hypothetical protein MGC3035	AL040896	Hs.22412	NP_077269
221985_at	0.010205	hypothetical protein FLJ20059	AW006750	Hs.246875	NP_060114
221992_at	0.042536	ESTs, Moderately similar to nuclear pore complex interacting protein [Homo sapiens] [H.sapiens]	AI925734	Hs.409134	
222023_at	0.018693	Consensus includes gb:AK022014.1 /DEF=Homo sapiens cDNA FLJ11952 fis, clone HEMBB1000831, weakly similar to Homo sapiens breast cancer nuclear receptor-binding auxiliary protein (BRX) mRNA. /FEA=mRNA /DB_XREF=gi:10433327 /UG=Hs.306619 Homo sapiens cDNA FLJ11952 fis, clone HEMBB1000831, weakly similar to Homo sapiens breast cancer nuclear receptor-binding auxiliary protein (BRX) mRNA	AK022014		NP_658913

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222029_x_at	0.025293	Consensus includes gb:NM_014260.1 /DEF=Homo sapiens HLA class II region expressed gene KE2 (HKE2), mRNA. /FEA=CDS /GEN=HKE2 /PROD=HLA class II region expressed gene KE2 /DB_XREF=gi:7657161 /UG=Hs.205736 HLA class II region expressed gene KE2 /FL=gb:NM_014260.1	NM_014260		NP_055075
222035_s_at	0.003171	poly(A) polymerase alpha	AI984479	Hs.49007	NP_116021
222036_s_at	0.0213	MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AI859865	Hs.154443	
222039_at	0.006443	ESTs, Moderately similar to PRO0478 protein [Homo sapiens] [H.sapiens]	AA292789	Hs.375584	
222040_at	0.013088	heterogeneous nuclear ribonucleoprotein A1	AI144007	Hs.249495	NP_112420
222041_at	0.021367	ESTs	BG235929	Hs.391830	NP_001374
222052_at	0.005852	ESTs, Weakly similar to 2109260A B cell growth factor [Homo sapiens] [H.sapiens]	AA001552	Hs.59839	
222061_at	0.004908	CD58 antigen, (lymphocyte function-associated antigen 3)	AA700015	Hs.75626	
222065_s_at	0.02145	flightless I homolog (Drosophila)	AI830227	Hs.83849	NP_002009
222067_x_at	0.012818	Consensus includes gb:AL353759 /DEF=Human DNA sequence from clone RP1-221C16 on chromosome 6. Contains two genes for novel histone 4 family members, two genes for novel histone 1 family members, three genes for novel histone 2B family members, a gene for a novel histone 2A family me... /FEA=mRNA_10 /DB_XREF=gi:8745068 /UG=Hs.180779 H2B histone family, member B	AL353759		
222071_s_at	1.26E-04	ESTs, Highly similar to hypothetical protein PRO2176 [Homo sapiens] [H.sapiens]	BE552428	Hs.391502	
222073_at	0.002418	collagen, type IV, alpha 3 (Goodpasture antigen)	AI694562	Hs.530	NP_112734
222077_s_at	7.58E-04	Rac GTPase activating protein 1	AU153848	Hs.23900	NP_037409

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222082_at	0.016934	HIV-1 inducer of short transcripts binding protein; lymphoma related factor	AI568395	Hs.104640	NP_056982
222111_at	0.00161	ESTs, Moderately similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AU145293	Hs.61773	
222119_s_at	1.26E-04	Consensus includes gb:AL117620.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564K2364 (from clone DKFZp564K2364); partial cds. /FEA=mRNA /GEN=DKFZp564K2364 /PROD=hypothetical protein /DB_XREF=gi:5912200 /UG=Hs.284289 vitiligo-associated protein VIT-1	AL117620		NP_079409
222129_at	0.02145	Consensus includes gb:AK026155.1 /DEF=Homo sapiens cDNA: FLJ22502 fis, clone HRC11383. /FEA=mRNA /DB_XREF=gi:10438915 /UG=Hs.298140 Homo sapiens cDNA: FLJ22502 fis, clone HRC11383	AK026155		NP_077269
222132_s_at	0.007052	Consensus includes gb:AJ278150.1 /DEF=Homo sapiens mRNA for putative lipid kinase. /FEA=mRNA /PROD=putative lipid kinase /DB_XREF=gi:8250242 /UG=Hs.260238 hypothetical protein FLJ10842	AJ278150		NP_060708
222133_s_at	0.002418	Consensus includes gb:AK022280.1 /DEF=Homo sapiens cDNA FLJ12218 fis, clone MAMMA1001075, moderately similar to Homo sapiens CGI-72 protein mRNA. /FEA=mRNA /DB_XREF=gi:10433640 /UG=Hs.288435 Homo sapiens cDNA FLJ12218 fis, clone MAMMA1001075, moderately similar to Homo sapiens CGI-72 protein mRNA	AK022280		NP_057102

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222143_s_at	0.001116	Consensus includes gb:AY007098.1 /DEF=Homo sapiens clone TCCCIA00183 mRNA sequence. /FEA=mRNA /DB_XREF=gi:9955989 /UG=Hs.27556 hypothetical protein FLJ22405	AY007098		NP_071930
222148_s_at	3.26E-04	hypothetical protein FLJ11040	BF688108	Hs.14202	NP_060777
222151_s_at	0.016934	Consensus includes gb:AK023738.1 /DEF=Homo sapiens cDNA FLJ13676 fis, clone PLACE1011922, weakly similar to MYOSIN HEAVY CHAIN, NONMUSCLE TYPE B. /FEA=mRNA /DB_XREF=gi:10435760 /UG=Hs.300876 hypothetical protein FLJ13386	AK023738		NP_079456
222156_x_at	1.26E-04	Consensus includes gb:AK022459.1 /DEF=Homo sapiens cDNA FLJ12397 fis, clone MAMMA1002769, weakly similar to Homo sapiens cell cycle progression restoration 8 protein (CPR8) mRNA. /FEA=mRNA /DB_XREF=gi:10433861 /UG=Hs.82506 KIAA1254 protein	AK022459		NP_004739
222158_s_at	0.002418	Consensus includes gb:AF229834.1 /DEF=Homo sapiens apoptosis-related protein PNAS-4 (PNAS-4) mRNA, partial cds. /FEA=mRNA /GEN=PNAS-4 /PROD=apoptosis-related protein PNAS-4 /DB_XREF=gi:7229639 /UG=Hs.42409 CGI-146 protein	AF229834		NP_057160
222165_x_at	0.04734	Consensus includes gb:AK022885.1 /DEF=Homo sapiens cDNA FLJ12823 fis, clone NT2RP2002752. /FEA=mRNA /DB_XREF=gi:10434538 /UG=Hs.95867 hypothetical protein EST00098	AK022885		NP_077017

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222173_s_at	0.001253	Consensus includes gb:AK026105.1 /DEF=Homo sapiens cDNA: FLJ22452 fis, clone HRC09667. /FEA=mRNA /DB_XREF=gi:10438847 /UG=Hs.135917 hypothetical protein DKFZp761D1823	AK026105		NP_060891
222175_s_at	0.04734	Consensus includes gb:AK000003.1 /DEF=Homo sapiens mRNA for FLJ00003 protein, partial cds. /FEA=mRNA /GEN=FLJ00003 /PROD=FLJ00003 protein /DB_XREF=gi:7209306 /UG=Hs.8657 TIG-1	AK000003		NP_056973
222186_at	0.0174	Consensus includes gb:AL109684.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 27080. /FEA=mRNA /DB_XREF=gi:5689805 /UG=Hs.306329 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 27080	AL109684		
222187_x_at	0.007934	Consensus includes gb:X78262.1 /DEF=H.sapiens mRNA for TRE5. /FEA=mRNA /DB_XREF=gi:587440 /UG=Hs.302178 H.sapiens mRNA for TRE5	X78262		
222204_s_at	0.021165	Consensus includes gb:AL110238.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566E104 (from clone DKFZp566E104); partial cds. /FEA=mRNA /GEN=DKFZp566E104 /PROD=hypothetical protein /DB_XREF=gi:5817180 /UG=Hs.189834 DKFZP566E104 protein	AL110238		NP_060897
222207_x_at	0.010205	Consensus includes gb:AK024602.1 /DEF=Homo sapiens cDNA: FLJ20949 fis, clone ADSE01902. /FEA=mRNA /DB_XREF=gi:10436915 /UG=Hs.220255 Homo sapiens cDNA: FLJ20949 fis, clone ADSE01902	AK024602		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222218_s_at	0.016934	Consensus includes gb:AJ400843.1 /DEF=Homo sapiens partial mRNA for immunoglobulin-like cell surface receptor FDF03-M14, soluble alternative form. /FEA=mRNA /GEN=FDF03-M14 /PROD=cell surface receptor FDF03-M14 /DB_XREF=gi:9715838 /UG=Hs.122591 paired immunoglobulin-like receptor alpha	AJ400843		NP_840057
222230_s_at	1.26E-04	Consensus includes gb:AK022248.1 /DEF=Homo sapiens cDNA FLJ12186 fis, clone MAMMA1000824, weakly similar to ACTIN. /FEA=mRNA /DB_XREF=gi:10433604 /UG=Hs.274369 uncharacterized hypothalamus protein HARP11	AK022248		
222231_s_at	1.26E-04	Consensus includes gb:AK025328.1 /DEF=Homo sapiens cDNA: FLJ21675 fis, clone COL09090, highly similar to AF119857 Homo sapiens PRO1855 mRNA. /FEA=mRNA /DB_XREF=gi:10437822 /UG=Hs.283558 hypothetical protein PRO1855	AK025328		NP_060979
222233_s_at	0.016934	Consensus includes gb:AK022922.1 /DEF=Homo sapiens cDNA FLJ12860 fis, clone NT2RP2003559. /FEA=mRNA /DB_XREF=gi:10434591 /UG=Hs.28891 hypothetical protein FLJ11360	AK022922		NP_071932
222235_s_at	2.05E-04	match: proteins: Tr:O02330 Tr:Q9Y2J5 Tr:Q9V5I7; Human DNA sequence from clone RP1-19N1 on chromosome Xq21.33-22.3 Contains a gene for a novel protein. Contains ESTs, STSs and GSSs, complete sequence.	AL139812		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222244_s_at	0.007779	Consensus includes gb:AK000749.1 /DEF=Homo sapiens cDNA FLJ20742 fis, clone HEP06891. /FEA=mRNA /DB_XREF=gi:7021031 /UG=Hs.52184 hypothetical protein FLJ20618	AK000749		NP_060373
222262_s_at	0.003368	Consensus includes gb:AL137750.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E1212 (from clone DKFZp434E1212). /FEA=mRNA /DB_XREF=gi:6808386 /UG=Hs.120439 ethanolamine kinase	AL137750		NP_061108
222266_at	6.68E-04	chromosome 19 open reading frame 2	BF796940	Hs.7943	NP_604431
222294_s_at	1.26E-04	ESTs	AW971415	Hs.165337	
222311_s_at	0.04734	ESTs, Highly similar to SRA4_HUMAN CTD-binding SR-like protein RA4 [H.sapiens]	AA648521	Hs.390734	
222326_at	0.003539	ESTs	AW973834	Hs.105884	
222366_at	3.26E-04	ESTs	W86781	Hs.293736	
222368_at	1.26E-04	hypothetical protein from EUROIMAGE 1759349	AW972351	Hs.107203	NP_660274
266_s_at	2.05E-04	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	NM_013230	Hs.381003	NP_037362
31799_at	0.005276	EST	AF070618	Hs.14074	
31826_at	0.026891	KIAA0674 protein	AB014574	Hs.14799	
31835_at	0.025464	histidine-rich glycoprotein precursor; Human histidine-rich glycoprotein mRNA, complete cds.	M13149		
31845_at	0.005852	E74-like factor 4 (ets domain transcription factor)	NM_001421	Hs.151139	NP_001412
31861_at	0.016934	immunoglobulin mu binding protein 2	NM_002180	Hs.1521	NP_002171
32029_at	0.012681	3-phosphoinositide dependent protein kinase-1	NM_002613	Hs.154729	NP_002604
32032_at	7.58E-04	EST, Weakly similar to T34549 probable mucin protein DKFZp434B0635.1 - human (fragment) [H.sapiens]	NM_022719	Hs.400995	NP_073210
32091_at	1.26E-04	KIAA0446 gene product	AB007915	Hs.158286	
32836_at	0.001618	1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)	U56417		NP_116130
33494_at	0.025245	electron-transferring-flavoprotein dehydrogenase	NM_004453	Hs.323468	NP_004444

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
34031_i_at	0.004341	cerebral cavernous malformations 1	NM_004912	Hs.93810	NP_004903
34206_at	0.016934	centaurin, delta 2	AB018325	Hs.21264	NP_631920
34210_at	2.05E-04	CDW52 antigen (CAMPATH-1 antigen)	N90866	Hs.276770	NP_001794
34408_at	2.05E-04	reticulon 2	NM_005619	Hs.3803	NP_005610
34858_at	0.041055	KIAA0176 protein	D79998	Hs.4935	
35254_at	7.58E-04	FLN29 gene product	NM_006700	Hs.5148	NP_006691
35776_at	0.048473	intersectin 1 (SH3 domain protein)	NM_003024	Hs.66392	NP_003015
36019_at	2.05E-04	dom-3 homolog Z (C. elegans)	L26260	Hs.444	NP_115830
36129_at	0.004341	KIAA0397 gene product	AB007857	Hs.7416	
36545_s_at	0.003171	KIAA0542 gene product	AB011114	Hs.62209	
36554_at	0.035611	Cluster Incl. Y15521:Homo sapiens ASMTL gene /cds=(0,1889) /gb=Y15521 /gi=3808147 /ug=Hs.6315 /len=1890	Y15521		
36564_at	0.016934	hypothetical protein FLJ90005	W27419	Hs.352358	
36566_at	0.028067	cystinosis, nephropathic	NM_004937	Hs.64837	NP_004928
36907_at	0.003171	mevalonate kinase (mevalonic aciduria)	NM_000431	Hs.130607	NP_000422
36920_at	0.027243	myotubular myopathy 1	NM_000252	Hs.75302	NP_000243
37012_at	0.003171	capping protein (actin filament) muscle Z-line, beta	NM_004930	Hs.333417	NP_004921
37145_at	0.033381	Homo sapiens NKG5 gene, complete cds.	M85276		
37170_at	0.003539	BMP-2 inducible kinase	NM_017593	Hs.198642	NP_060063
37232_at	0.004341	translocase of inner mitochondrial membrane 9 homolog (yeast)	NM_014749	Hs.77724	NP_055564
37278_at	0.007779	tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome)	NM_000116	Hs.79021	NP_851831
37943_at	0.00161	KIAA0321 protein	AB002319	Hs.8663	
37965_at	0.010205	parvin, beta	AA181053	Hs.8836	NP_037459
37966_at	0.005276	parvin, beta	AA187563	Hs.8836	NP_037459
37986_at	0.009179	erythropoietin receptor	NM_000121	Hs.127826	NP_000112
38241_at	0.010205	butyrophilin, subfamily 3, member A2	NM_006994	Hs.167741	NP_008925
38671_at	0.004341	KIAA0620 protein	AB014520	Hs.301685	NP_055918
38892_at	0.010205	KIAA0240 protein	D87077	Hs.196275	
38964_r_at	0.007779	Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	NM_000377	Hs.2157	NP_000368
39313_at	0.016934	protein kinase, lysine deficient 1	NM_018979	Hs.184592	NP_061852
39318_at	0.018693	T-cell leukemia/lymphoma 1A	NM_021966	Hs.2484	NP_068801
396_f_at	0.020793	erythropoietin receptor	NM_000121	Hs.127826	NP_000112
39650_s_at	7.58E-04	KIAA0435 gene product	NM_014801	Hs.31438	NP_055616
39729_at	5.03E-04	Human natural killer cell enhancing factor (NKEFB) mRNA, complete cds.	NM_005809		NP_005800
39817_s_at	0.013223	putative c-Myc-responsive	NM_006443	Hs.109752	NP_006434

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
39891_at	3.26E-04	Homo sapiens cDNA FLJ37393 fis, clone BRAMY2027073, moderately similar to Fugu rubripes zinc finger protein	AI246730	Hs.126901	
40016_g_at	0.021876	KIAA0303 protein	AB002301	Hs.54985	
40020_at	0.021876	cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila)	NM_001407	Hs.55173	NP_001398
40149_at	7.58E-04	SH2-B homolog	NM_015503	Hs.15744	NP_056318
40225_at	0.026891	cyclin G associated kinase	NM_005255	Hs.153227	NP_005246
40255_at	0.007779	Cluster Incl. AC004531:Homo sapiens Chromosome 16 BAC clone CIT987SK-A-67A1 /cds=(0,1622) /gb=AC004531 /gi=3337392 /ug=Hs.155049 /len=1623	AC004531		
41047_at	0.026891	chromosome 9 open reading frame 16	AI885170	Hs.95867	NP_077017
41469_at	0.003171	elafin has been sequenced at the protein level; pre-elafin has not; its existence is assumed from its molecular weight (PAGE analysis); putative; Homo sapiens elafin precursor, gene, complete cds.	NM_002638		NP_002629
41512_at	0.026891	BRCA1 associated protein	AL042733	Hs.122764	NP_006759
41856_at	0.013223	EST	AL049370	Hs.13350	
44111_at	0.00161	vacuolar protein sorting 33B (yeast)	AI672363	Hs.26510	NP_061138
44696_at	5.03E-04	hypothetical protein FLJ10743	AA915989	Hs.3376	NP_060671
44790_s_at	0.041055	hypothetical protein FLJ21562	AI129310	Hs.288708	NP_079389
45526_g_at	3.26E-04	hypothetical protein FLJ14154	AI246641	Hs.184870	NP_079121
45828_at	5.03E-04	hypothetical protein FLJ10241	AI768100	Hs.299867	NP_060505
46270_at	2.05E-04	ubiquitin associated protein 1	AL039447	Hs.75425	NP_057609
46323_at	7.58E-04	Ca ²⁺ -dependent endoplasmic reticulum nucleoside diphosphatase	AL120741	Hs.389537	NP_620148
47069_at	0.00228	Rho GTPase activating protein 8	AA533284	Hs.102336	NP_851852
47083_at	0.02145	hypothetical protein MGC2718	AI280108	Hs.81057	NP_076972
47105_at	0.043705	hypothetical protein FLJ20399	AA886893	Hs.8575	NP_060273
47530_at	0.016934	hypothetical protein HSPC219	AA748492	Hs.9196	NP_057565
48106_at	0.010205	hypothetical protein FLJ20489	H14241	Hs.306989	NP_060312
48531_at	0.026891	TNFAIP3 interacting protein 2	AA522816	Hs.288991	NP_077285
48612_at	0.00161	hypothetical protein FLJ31821	AA225490	Hs.351575	NP_694574
48808_at	0.014715	dihydrofolate reductase	AI144299	Hs.83765	NP_000782
48825_at	0.007934	candidate tumor suppressor p33 ING1 homolog	AA887083	Hs.108183	NP_057246
49077_at	0.00228	protein phosphatase methylesterase-1	AL040538	Hs.63304	NP_057231
49306_at	0.003539	AD037 protein	AI890191	Hs.296162	NP_835281

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
49327_at	0.033381	sirtuin silent mating type information regulation 2 homolog 3 (S. cerevisiae)	AI492888	Hs.19306	NP_036371
49679_s_at	0.011862	matrix metalloproteinase 24 (membrane-inserted)	AA243774	Hs.355907	NP_006681
50314_i_at	0.005039	chromosome 20 open reading frame 27	AI761506	Hs.274422	NP_060344
51146_at	2.17E-04	hypothetical protein FLJ20477	AA203365	Hs.26994	
51176_at	0.004341	cofactor required for Sp1 transcriptional activation, subunit 8, 34kDa	AA131335	Hs.28166	NP_004260
51200_at	0.013223	hypothetical protein FLJ20850	AI744084	Hs.30783	NP_060437
51774_s_at	0.041055	ESTs, Moderately similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW014299	Hs.237946	
52159_at	0.019761	HEMK homolog 7kb	W93807	Hs.46907	NP_057257
52169_at	0.00161	hypothetical protein FLJ90524	AI302185	Hs.388506	NP_699166
53071_s_at	0.041055	hypothetical protein FLJ22222	AI885411	Hs.54558	NP_787098
53912_at	7.58E-04	sorting nexin 11	AI668643	Hs.15827	NP_689450
55081_at	0.02145	chromosome 22 open reading frame 23	W46406	Hs.8535	NP_203744
55692_at	2.05E-04	engulfment and cell motility 2 (ced-12 homolog, C. elegans)	W22924	Hs.96560	NP_573403
55872_at	0.028893	KIAA1196 protein	AI493119	Hs.158654	
57588_at	0.046329	solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	R62432	Hs.12321	
57703_at	0.035171	hypothetical protein MGC27076	N32782	Hs.13138	NP_689912
57715_at	0.026891	hypothetical protein LOC51063	W72694	Hs.241545	NP_057000
59644_at	7.58E-04	BMP-2 inducible kinase	AI735391	Hs.198642	NP_060063
59999_at	0.016934	hypoxia-inducible factor 1, alpha subunit inhibitor	W37897	Hs.14595	NP_060372
60528_at	0.033381	phospholipase A2, group IVB (cytosolic)	N71116	Hs.198161	NP_005081
60794_f_at	3.26E-04	Human zinc finger protein zfp6 (ZF6) mRNA, partial cds	AI400621	Hs.334690	
62212_at	0.042536	hypothetical protein MGC955	W37846	Hs.55608	NP_077002
632_at	0.010205	glycogen synthase kinase 3 alpha	NM_019884	Hs.118890	NP_063937
64064_at	0.007779	immune associated nucleotide 4 like 1 (mouse)	AI435089	Hs.26194	NP_060854
64371_at	0.033491	KIAA0365 gene product	AI978718	Hs.190452	
64408_s_at	3.09E-04	serologically defined breast cancer antigen NY-BR-20	AW025529	Hs.239812	
64438_at	0.00228	hypothetical protein FLJ22222	W19668	Hs.54558	NP_787098
64440_at	0.034221	interleukin 17 receptor C	AI560217	Hs.129959	NP_703193
64486_at	7.58E-04	coronin, actin-binding protein, 1B	AI341234	Hs.408995	NP_065174
64488_at	0.019879	Homo sapiens FKSG27 (FKSG27) mRNA, complete cds	AW003091	Hs.6217	
64883_at	1.26E-04	hypothetical protein MGC26706	AI744083	Hs.65406	NP_689794

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
65630_at	0.033381	Homo sapiens, Similar to RIKEN cDNA 5530601I19 gene, clone MGC:9743 IMAGE:3854028, mRNA, complete cds	AI742455	Hs.28974	NP_060739
74694_s_at	0.005852	hypothetical protein FLJ23282	AA907940	Hs.170253	NP_079092
78047_s_at	0.010205	matrix metalloproteinase 24 (membrane-inserted)	AW001777	Hs.355907	NP_006681
78383_at	5.03E-04	tumor protein p53-binding protein	AI150117	Hs.179982	NP_005793
89476_r_at	0.003928	aminopeptidase-like 1	AA398062	Hs.71746	NP_078939
90610_at	0.041055	leucine-rich repeat protein, neuronal 1	AI654857	Hs.125742	NP_002310
91684_g_at	0.01185	exosome component Rrp41	AI571298	Hs.343589	NP_061910
AFFX-BioB-3_at	0.00161	J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423		
AFFX-BioC-3_at	0.013223	J04423 E coli bioC protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-BioDn-5_at	0.004341	J04423 E coli bioD gene dethiobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-hum_alu_at	0.003171	***ALU WARNING: Human Alu-Sq subfamily consensus sequence.	U14573		
AFFX-HUMGAPDH/M33197_3_at	0.001116	glyceraldehyde-3-phosphate dehydrogenase	NM_002046	Hs.169476	NP_002037
AFFX-HUMRGE/M10098_3_at	0.003539	Human 18S rRNA gene, complete.	M10098		
AFFX-HUMRGE/M10098_M_at	6.68E-04	Human 18S rRNA gene, complete.	M10098		
AFFX-M27830_5_at	1.26E-04	Human 28S ribosomal RNA gene, complete cds.	M27830		
AFFX-r2-Ec-bioB-3_at	0.003171	Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2772-3004 of J04423 /LEN=1114 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
AFFX-r2-Ec-bioC-3_at	0.033381	Escherichia coli /REF=J04423 /DEF=E coli bioC protein corresponding to nucleotides 4609-4883 of J04423 /LEN=777 (5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-r2-Ec-bioD-5_at	0.003171	Escherichia coli /REF=J04423 /DEF=E coli bioD gene dethiobiotin synthetase corresponding to nucleotides 5024-5244 of J04423 /LEN=676 (5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-r2-Hs18SrRNA-3_s_at	0.003539	Human 18S rRNA gene, complete.	M10098		
AFFX-r2-Hs18SrRNA-M_x_at	0.018693	Human 18S rRNA gene, complete.	M10098		
AFFX-r2-Hs28SrRNA-3_at	1.26E-04	Human 28S ribosomal RNA gene.	M11167		
AFFX-r2-P1-cre-3_at	0.02145	Bacteriophage /REF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 1032-1270 of X03453 /LEN=1058 (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	X03453		